## SEQUENCE LISTING

```
<110> INCYTE PHARMACEUTICALS, INC.
       HILLMAN, Jennifer L.
BANDMAN, Olga
       LAL, Preeti
       YUE, Henry
       REDDY, Roopa
       TANG, Y. Tom
       GERSTIN, Edward H.
       PATTERSON, Chandra
      BAUGHN, Mariah R.
      AZIMZAI, Yalda
      LU, Dyung Aina M.
<120> HUMAN TRANSCRIPTIONAL REGULATOR MOLECULES
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<141> Herewith
<150> 60/084,254; 60/095,827; 60/102,745
<151> 1998-05-05; 1998-08-07; 1998-10-02
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Ser Cys Phe Leu Arg Leu Gly Arg Ser Thr Leu Leu Glu Leu Glu
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Pro Ala Gly Arg Pro Cys Ser Gly Arg Thr Arg His Arg Ala Leu
                  35
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His Arg Arg Leu Val Ala Cys Val Thr Val Ser Ser Arg Arg His
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Arg Lys Glu Ala Gly Arg Gly Arg Ala Glu Ser Phe Ile Ala Val
                 65
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Gly Met Ala Ala Pro Ser Met Lys Glu Arg Gln Val Cys Trp Gly
                 80
                                      85
                                                           90
Ala Arg Asp Glu Tyr Trp Lys Cys Leu Asp Glu Asn Leu Glu Asp
                 95
                                     100
                                                           105
Ala Ser Gln Cys Lys Leu Arg Ser Ser Phe Glu Ser Ser Cys
                110
                                     115
                                                          120
Pro Gln Gln Trp Ile Lys Tyr Phe Asp Lys Arg Arg Asp Tyr Leu
                 125
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Lys Phe Lys Glu Lys Phe Glu Ala Gly Gln Phe Glu Pro Ser Glu
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Thr Thr Ala Lys Ser
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Leu Leu Val Tyr Asp Met Asn Leu Arg Glu Met Glu Asn Tyr Glu
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Lys Ile Tyr Lys Glu Ile Glu Cys Ser Ile Ala Gly Ala His Glu
                 35
                                      40
                                                           45
Lys Ile Ala Glu Cys Lys Lys Gln Ile Leu Gln Ala Lys Arg Ile
                 50
                                      55
Arg Lys Asn Arg Gln Glu Tyr Asp Ala Leu Ala Lys Val Ile Gln
                 65
                                      70
His His Pro Asp Arg His Glu Thr Leu Lys Glu Leu Glu Ala Leu
                 80
                                      85
Gly Lys Glu Leu Glu His Leu Ser His Ile Lys Glu Ser Val Glu
                 95
                                     100
Asp Lys Leu Glu Leu Arg Arg Lys Gln Phe His Val Leu Leu Ser
                110
                                     115
                                                          120
Thr Ile His Glu Leu Gln Gln Thr Leu Glu Asn Asp Glu Lys Leu
                125
                                     130
Ser Glu Val Glu Glu Ala Gln Glu Ala Ser Met Glu Thr Asp Pro
                140
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Lys Pro
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Ala Ala His Arg Phe Phe Cys His Phe Cys Lys Gly Glu Val Ser
                 20
                                      25
Pro Lys Leu Pro Glu Tyr Ile Cys Pro Arg Cys Glu Ser Gly Phe
                 35
                                      40
                                                          45
Ile Glu Glu Val Thr Asp Asp Ser Ser Phe Leu Gly Gly Gly
                 50
                                      55
Ser Arg Ile Asp Asn Thr Thr Thr Thr His Phe Ala Glu Leu Trp
                 65
                                      70
Gly His Leu Asp His Thr Met Phe Phe Gln Asp Phe Arg Pro Phe
                 80
                                      85
Leu Ser Ser Pro Leu Asp Gln Asp Asn Arg Ala Asn Glu Arg
                                     100
                                                         105
Gly His Gln Thr His Thr Asp Phe Trp Gly Ala Arg Pro Pro Arg
                110
                                     115
                                                         120
Leu Pro Leu Gly Arg Arg Tyr Arg Ser Arg Gly Ser Ser Arg Pro
                125
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Asp Arg Ser Pro Ala Ile Glu Gly Ile Leu Gln His Ile Phe Ala
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140
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Gly Phe Phe Ala Asn Ser Ala Ile Pro Gly Ser Pro His Pro Phe
                 155
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Ser Trp Ser Gly Met Leu His Ser Asn Pro Gly Asp Tyr Ala Trp
                170
                                     175
                                                          180
Gly Gln Thr Gly Leu Asp Ala Ile Val Thr Gln Leu Leu Gly Gln
                185
                                     190
                                                          195
Leu Glu Asn Thr Gly Pro Pro Pro Ala Asp Lys Glu Lys Ile Thr
                200
                                     205
                                                          210
Ser Leu Pro Thr Val Thr Val Thr Gln Glu Gln Val Asp Met Gly
                215
                                     220
                                                          225
Leu Glu Cys Pro Val Cys Lys Glu Asp Tyr Thr Val Glu Glu Glu
                230
                                     235
Val Arg Gln Leu Pro Cys Asn His Phe Phe His Ser Ser Cys Ile
                245
                                     250
                                                         255
Val Pro Trp Leu Glu Leu His Asp Thr Cys Pro Val Cys Arg Lys
                260
                                     265
Ser Leu Asn Gly Glu Asp Ser Thr Arg Gln Ser Gln Ser Thr Glu
                275
                                    280
                                                         285
Ala Ser Ala Ser Asn Arg Phe Ser Asn Asp Ser Gln Leu His Asp
                                    295
Arg Trp Thr Phe
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Ser Val Asn Arg Lys Arg Leu Asn Arg Asn Ala Arg Arg Lys Ala
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                                      25
Ala Pro Arg Ile Glu Cys Ser His Ile Arg His Ala Trp Asp His
                 35
                                      40
Ala Lys Ser Val Arg Gln Asn Leu Ala Glu Met Gly Leu Ala Val
                 50
                                      55
Asp Pro Asn Arg Ala Val Pro Leu Arg Lys Arg Lys Val Lys Ala
                 65
                                      70
Met Glu Val Asp Ile Glu Glu Arg Pro Lys Glu Leu Val Arg Lys
                 80
                                      85
Pro Tyr Val Leu Asn Asp Leu Glu Ala Glu Ala Ser Leu Pro Glu
                 95
                                     100
Lys Lys Gly Asn Thr Leu Ser Arg Asp Leu Ile Asp Tyr Val Arg
                110
                                     115
Tyr Met Val Glu Asn His Gly Glu Asp Tyr Lys Ala Met Ala Arg
                125
                                     130
                                                          135
Asp Glu Lys Asn Tyr Tyr Gln Asp Thr Pro Lys Gln Ile Arg Ser
                140
                                     145
Lys Ile Asn Val Tyr Lys Arg Phe Tyr Pro Ala Glu Trp Gln Asp
                155
                                     160
Phe Leu Asp Ser Leu Gln Lys Arg Lys Met Glu Val Glu
                170
                                     175
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Trp Pro Gln Lys Pro Gln Cys His Gly Ser Gly Val Ile His Gly
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Asn Ser Pro Leu Cys Pro Asn Trp Gln Val Phe Pro Leu Val Arg
                 35
                                                           45
Pro His Arg Gln Ser Arg Gln Leu Gln Val Pro Glu Pro Ile Gln
                 50
                                      55
Ala Gly Gly Pro Ser Cys Gly His His Ser Pro Trp Arg Leu Phe
                 65
                                      70
Leu Pro Gln Arg Lys Ser Gln Val Ser Arg Gly Gly Arg Leu Ala
                 80
                                      85
Cys Leu Leu Ser Tyr Ala Gly Leu Ser Gly Asp Asp Pro Asp Leu
                 95
                                     100
Gly Pro Ala His Val Val Thr Val Ile Ala Arg Gln Arg Gly Asp
                110
                                     115
                                                          120
Gln Leu Val Pro Phe Ser Thr Lys Ser Gly Asp Thr Leu Leu
                125
                                     130
                                                          135
Leu His His Gly Asp Phe Ser Ala Glu Glu Val Phe His Arg Glu
                140
                                     145
                                                         150
Leu Arg Ser Asn Ser Met Lys Thr Trp Gly Leu Arg Ala Ala Gly
                155
                                     160
Trp Met Ala Met Phe Met Gly Leu Asn Leu Met Thr Arg Ile Leu
                170
                                     175
                                                         180
Tyr Thr Leu Val Asp Trp Phe Pro Val Phe Arg Asp Leu Val Asn
                185
                                     190
Ile Gly Leu Lys Ala Phe Ala Phe Cys Val Ala Thr Ser Leu Thr
                200
                                     205
                                                         210
Leu Leu Thr Val Ala Ala Gly Trp Leu Phe Tyr Arg Pro Leu Trp
                215
                                     220
Ala Leu Leu Ile Ala Gly Leu Ala Leu Val Pro Ile Leu Val Ala
                230
                                     235
Arg Thr Arg Val Pro Ala Lys Lys Leu Glu
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Leu Leu Glu Glu Ser Gly Asp Leu Gly Thr Ala Pro Asp Glu Ala
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                                      25
Val Arg Ala Pro Leu Asp Trp Ala Leu Pro Leu Ser Glu Val Pro
                 35
                                      40
Ser Asp Trp Glu Val Asp Asp Leu Leu Cys Ser Leu Leu Ser Pro
                                      55
Pro Ala Ser Leu Asn Ile Leu Ser Ser Ser Asn Pro Cys Leu Val
                 65
                                                          75
His His Asp His Thr Tyr Ser Leu Pro Arg Glu Thr Val Ser Met
                 80
                                      85
Asp Leu Glu Ser Glu Ser Cys Arg Lys Glu Gly Thr Gln Met Thr
                 95
                                     100
                                                         105
Pro Gln His Met Glu Glu Leu Ala Glu Gln Glu Ile Ala Arg Leu
                110
                                     115
Val Leu Thr Asp Glu Glu Lys Ser Leu Leu Glu Lys Glu Gly Leu
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<210> 7 <211> 371

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Ile	Leu	Pro	Glu	Thr 140	Leu	Pro	Leu	Thr		Thr	Glu	Glu	Gln	Ile 150
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				170				Tyr	175		_			Ser 180
				185				Asn	190					Lys 195
				200				Leu	205					210
				215				Glu	220					225
				230				Leu	235					240
				245				Ser	250					255
				260				Arg	265					270
				275				Leu	280					285
				290				Trp	295					300
				305				Ser	310					315
				320				Pro	325					330
				335				Arg	340					345
				350				Gly	355		Pro	Thr	Gly	Ser 360
Pro	Ser	Val	Ile	Leu 365	Gln	Asp	Arg	Tyr	Ser 370	Gly				

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Asp Gly Asp Glu Asp Arg Asn Pro Ser Thr Ala Phe Tyr Gln Ala
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                                     25
                                                          30
Phe His Leu Asn Thr Leu Lys Glu Ser Lys Ser Leu Trp Asp Ser
                 35
                                     40
Ala Ser Gly Gly Val Val Ala Ile Asp Asn Lys Ile Glu Gln
                 50
                                     55
Ala Met Asp Leu Val Lys Ser His Leu Met Tyr Ala Val Arg Glu
                 65
                                     70
                                                         75
Glu Val Glu Val Leu Lys Glu Gln Ile Lys Glu Leu Val Glu Arg
                 80
                                     85
Asn Ser Leu Leu Glu Arg Glu Asn Ala Leu Leu Lys Ser Leu Ser
                 95
                                    100
Ser Asn Asp Gln Leu Ser Gln Leu Pro Thr Gln Gln Ala Asn Pro
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                                    115
Gly Ser Thr Ser Gln Gln Gln Ala Val Ile Ala Gln Pro Pro Gln
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<210> 8 <211> 148

Pro Thr Gln Pro Pro Gln Gln Pro Asn Val Ser Ser Ala
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135

<210> 9 <211> 127 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte clone 1760185CD1

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Val Leu Asn Lys Ala Ile Lys Ile Thr Val Pro Lys Gln Ser Asp
                110
                                     115
Ser Gln Pro Asn Ser Asp Asn Leu Ser Arg His Gly Glu Cys Gly
                125
                                     130
                                                          135
Lys Lys Gln Val Ser Tyr Arg Thr Asp Ile Val Gly Gly Val Pro
                140
                                                          150
Ile Ile Thr Pro Thr Gln Lys Glu Glu Val Asn Glu Cys Gly Glu
                155
                                     160
Ser Ile Asp Arg Asn Asn Leu Lys Arg Ser Gln Ser His Leu Pro
                170
                                     175
Tyr Phe Thr Pro Lys Pro Pro Gln Asp Ser Ala Val Ile Lys Ala
                185
                                     190
Gly Tyr Cys Val Lys Gln Gly Ala Val Met Lys Asn Trp Lys Arg
                200
                                     205
                                                          210
Arg Tyr Phe Gln Leu Asp Glu Asn Thr Ile Gly Tyr Phe Lys Ser
                215
                                     220
                                                          225
Glu Leu Glu Lys Glu Pro Leu Arg Val Ile Pro Leu Lys Glu Val
                230
                                     235
                                                          240
His Lys Val Gln Glu Cys Lys Gln Ser Asp Ile Met Met Arg Asp
                245
                                     250
                                                          255
Asn Leu Phe Glu Ile Val Thr Thr Ser Arg Thr Phe Tyr Val Gln
                260
                                     265
Ala Asp Ser Pro Glu Glu Met His Ser Trp Ile Lys Ala Val Ser
                275
                                     280
                                                          285
Gly Ala Ile Val Ala Gln Arg Gly Pro Gly Arg Ser Ala Ser Ser
                290
                                     295
Met Arg Gln Ala Arg Arg Leu Ser Asn Pro Cys Ile Gln Arg Ser
                305
                                     310
                                                          315
Ile Pro Pro Val Leu Gln Asn Pro Asn Thr Leu Ser Val Leu Pro
                320
                                     325
Thr Gln Pro Pro Pro Pro His Ile Pro Gln Pro Leu Ala Ala Thr
                335
                                     340
Leu Trp Ser Gln Pro Leu Pro Trp Arg Ser Glu Asp Phe Thr Ser
                350
                                     355
Leu Leu Pro Arg Ser Ser Gln Gly Thr Ser Arg Ser Arg Leu Ser
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Leu Gln Glu Asn Gln Leu Pro Lys
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<223> Incyte clone 1850120CD1

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Met Pro Pro Pro Ser Glu Ile Gln Arg Leu Tyr Glu Leu Leu Ala
                110
                                    115
Ala His Gly Thr Leu Glu Leu Gln Ala Glu Ile Leu Pro Arg Arg
                125
                                    130
Pro Pro Thr Pro Glu Arg Gln Ser Glu Glu Glu Arg Ser Asp Glu
                140
                                    145
                                                         150
Glu Pro Glu Ala Lys Glu Glu Glu Glu Lys Pro His Met Pro
                155
                                    160
Thr Glu Phe Asp Phe Asp Asp Glu Pro Val Thr Pro Lys Asp Ser
                170
                                    175
Leu Ile Asp Arg Arg Thr Pro Gly Ser Ser Ala Arg Ser Gln
                185
                                    190
Lys Arg Glu Ala Arg Leu Asp Lys Val Leu Ser Asp Met Lys Arg
                200
                                    205
His Lys Lys Leu Glu Glu Gln Ile Leu Arg Thr Gly Arg Asp Leu
                215
                                    220
Phe Ser Leu Asp Ser Glu Asp Pro Ser Pro Ala Ser Pro Pro Leu
                230
                                    235
Arg Ser Ser Gly Ser Ser Leu Phe Pro Arg Gln Arg Lys Tyr
                245
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Ala Pro Pro Thr Val Ala Ala Pro Ala Pro Ser Leu Phe Pro Ala
                 20
                                      25
Ala Gln Met Met Asn Asn Gly Leu Leu Gln Gln Pro Ser Ala Leu
                  35
                                      40
                                                           45
Met Leu Leu Pro Cys Arg Pro Val Leu Thr Ser Val Ala Leu Asn
                 50
                                      55
Ala Asn Phe Val Ser Trp Lys Ser Arg Thr Lys Tyr Thr Ile Thr
                 65
                                      70
Pro Val Lys Met Arg Lys Ser Gly Gly Arg Asp His Thr Gly Arg
                 80
                                      85
Ile Arg Val His Gly Ile Gly Gly Gly His Lys Gln Arg Tyr Arg
                 95
                                     100
                                                          105
Met Ile Asp Phe Leu Arg Phe Arg Pro Glu Glu Thr Lys Ser Gly
                110
                                     115
                                                          120
Pro Phe Glu Glu Lys Val Ile Gln Val Arg Tyr Asp Pro Cys Arg
                125
                                     130
                                                          135
Ser Ala Asp Ile Ala Leu Val Ala Gly Gly Ser Arg Lys Arg Trp
                140
                                     145
                                                          150
Ile Ile Ala Thr Glu Asn Met Gln Ala Gly Asp Thr Ile Leu Asn
                155
                                     160
Ser Asn His Ile Gly Arg Met Ala Val Ala Ala Arg Glu Gly Asp
                170
                                     175
                                                          180
Ala His Pro Leu Gly Ala Leu Pro Val Gly Thr Leu Ile Asn Asn
                185
                                     190
                                                          195
Val Glu Ser Glu Pro Gly Arg Gly Ala Gln Tyr Ile Arg Ala Ala
                200
                                     205
                                                          210
Gly Thr Cys Gly Val Leu Leu Arg Lys Val Asn Gly Thr Ala Ile
                215
                                     220
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Ile Gln Leu Pro Ser Lys Arg Gln Met Gln Val Leu Glu Thr Cys
                230
                                     235
Val Ala Thr Val Gly Arg Val Ser Asn Val Asp His Asn Lys Arg
                245
                                     250
                                                          255
Val Ile Gly Lys Ala Gly Arg Asn Arg Trp Leu Gly Lys Arg Pro
                260
                                                          270
Asn Ser Gly Arg Trp His Arg Lys Gly Gly Trp Ala Gly Arg Lys
                275
                                     280
                                                          285
Ile Arg Pro Leu Pro Pro Met Lys Ser Tyr Val Lys Leu Pro Ser
                290
Ala Ser Ala Gln Ser
                305
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Leu Pro Glu Lys Val Ala Lys His Val Thr Leu Val Arg Glu Ser
                 20
                                      25
Gly Ser Leu Thr Tyr Glu Glu Phe Leu Gly Arg Val Ala Glu Leu
                                      40
Asn Asp Val Thr Ala Lys Val Ala Ser Gly Gln Glu Lys His Leu
                 50
Leu Phe Glu Val Gln Pro Gly Ser Asp Ser Ser Ala Phe Trp Lys
                 65
                                      70
                                                           75
Val Val Val Arg Val Val Cys Thr Lys Ile Asn Lys Ser Ser Gly
                 80
                                      85
Ile Val Glu Ala Ser Arg Ile Met Asn Leu Tyr Gln Phe Ile Gln
                 95
                                     100
                                                          105
Leu Tyr Lys Asp Ile Thr Ser Gln Ala Ala Gly Val Leu Ala Gln
                110
                                                          120
Ser Ser Thr Ser Glu Glu Pro Asp Glu Asn Ser Ser Ser Val Thr
                125
                                     130
                                                          135
Ser Cys Gln Ala Ser Leu Trp Met Gly Arg Val Lys Gln Leu Thr
                140
                                     145
Asp Glu Glu Glu Cys Cys Ile Cys Met Asp Gly Arg Ala Asp Leu
                155
                                     160
Ile Leu Pro Cys Ala His Ser Phe Cys Gln Lys Cys Ile Asp Lys
                170
                                     175
                                                          180
Trp Ser Asp Arg His Arg Asn Cys Pro Ile Cys Arg Leu Gln Met
                185
                                     190
Thr Gly Ala Asn Glu Ser Trp Val Val Ser Asp Ala Pro Thr Glu
                200
                                     205
                                                          210
Asp Asp Met Ala Asn Tyr Ile Leu Asn Met Ala Asp Glu Ala Gly
                215
                                     220
                                                          225
Gln Pro His Arg Pro
                230
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<221> misc feature <223> Incyte clone 2019742CB1

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<210> 15 <211> 232 <212> PRT <213> Homo sapiens

<220>

<221> misc\_feature

<223> Incyte clone 2056042CD1

290

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Glu Met Glu Glu Leu Arg Tyr Ala Pro Leu Ser Phe Arg Asn
                                     70
Pro Met Met Ser Lys Leu Arg Asn Tyr Arg Lys Asp Leu Ala Lys
                 80
                                     85
                                                          90
Leu His Arg Glu Val Arg Ser Thr Pro Leu Thr Ala Thr Pro Gly
                 95
                                     100
                                                         105
Gly Arg Gly Asp Met Lys Tyr Gly Ile Tyr Ala Val Glu Asn Glu
                110
                                     115
                                                         120
His Met Asn Arg Leu Gln Ser Gln Arg Ala Met Leu Leu Gln Gly
                125
                                     130
                                                         135
Thr Glu Ser Leu Asn Arg Ala Thr Gln Ser Ile Glu Arg Ser His
                140
                                     145
Arg Ile Ala Thr Glu Thr Asp Gln Ile Gly Ser Glu Ile Ile Glu
                155
                                    160
Glu Leu Gly Glu Gln Arg Asp Gln Leu Glu Arg Thr Lys Ser Arg
                170
                                    175
                                                         180
Leu Val Asn Thr Ser Glu Asn Leu Ser Lys Ser Arg Lys Ile Leu
                185
                                    190
                                                         195
Arg Ser Met Ser Arg Lys Val Thr Thr Asn Lys Leu Leu Ser
                200
                                    205
                                                         210
Ile Ile Ile Leu Leu Glu Leu Ala Ile Leu Gly Gly Leu Val Tyr
                215
                                    220
Tyr Lys Phe Phe Arg Ser His
                230
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Met Arg Gly Lys Thr Phe Arg Phe Glu Met Gln Arg Asp Leu Val
 1
                                      7.0
                                                           15
Ser Phe Pro Leu Ser Pro Ala Val Arg Val Lys Leu Val Ser Ala
                 20
Gly Phe Gln Thr Ala Glu Glu Leu Leu Glu Val Lys Pro Ser Glu
                 35
                                      40
Leu Ser Lys Glu Val Gly Ile Ser Lys Ala Glu Ala Leu Glu Thr
                 50
                                      55
Leu Gln Ile Ile Arg Arg Glu Cys Leu Thr Asn Lys Pro Arg Tyr
                 65
                                      70
Ala Gly Thr Ser Glu Ser His Lys Lys Cys Thr Ala Leu Glu Leu
                 80
                                      85
Leu Glu Gln Glu His Thr Gln Gly Phe Ile Ile Thr Phe Cys Ser
                 95
                                     100
                                                          105
Ala Leu Asp Asp Ile Leu Gly Gly Gly Val Pro Leu Met Lys Thr
                110
                                     115
                                                          120
Thr Glu Ile Cys Gly Ala Pro Gly Val Gly Lys Thr Gln Leu Cys
                125
                                     130
                                                          135
Met Gln Leu Ala Val Asp Val Gln Ile Pro Glu Cys Phe Gly Gly
                                     145
                                                          150
Val Ala Gly Glu Ala Val Phe Ile Asp Thr Glu Gly Ser Phe Met
                155
                                     160
Val Asp Arg Val Val Asp Leu Ala Thr Ala Cys Iie Gln His Leu
                170
                                     175
                                                          180
Gln Leu Ile Ala Glu Lys His Lys Gly Glu Glu His Arg Lys Ala
                185
                                     190
```

```
Leu Glu Asp Phe Thr Leu Asp Asn Ile Leu Ser His Ile Tyr Tyr
                 200
                                     205
                                                          210
 Phe Arg Cys Arg Asp Tyr Thr Glu Leu Leu Ala Gln Val Tyr Leu
                                     220
 Leu Pro Asp Phe Leu Ser Glu His Ser Lys Val Arg Leu Val Ile
                 230
                                                          240
 Val Asp Gly Ile Ala Phe Pro Phe Arg His Asp Leu Asp Asp Leu
                 245
                                     250
Ser Leu Arg Thr Arg Leu Leu Asn Gly Leu Ala Gln Gln Met Ile
                 260
                                     265
Ser Leu Ala Asn Asn His Arg Leu Ala Val Ile Leu Thr Asn Gln
                 275
                                     280
Met Thr Thr Lys Ile Asp Arg Asn Gln Ala Leu Leu Val Pro Ala
                 290
                                     295
Leu Gly Glu Ser Trp Gly His Ala Ala Thr Ile Arg Leu Ile Phe
                 305
                                     310
His Trp Asp Arg Lys Gln Arg Leu Ala Thr Leu Tyr Lys Ser Pro
                 320
                                     325
Ser Gln Lys Glu Cys Thr Val Leu Phe Gln Ile Lys Pro Gln Gly
                 335
                                     340
                                                          345
Phe Arg Asp Thr Val Val Thr Ser Ala Cys Ser Leu Gln Thr Glu
                 350
                                     355
Gly Ser Leu Ser Thr Arg Lys Arg Ser Arg Asp Pro Glu Glu Glu
                 365
                                     370
Leu
<210> 17
<211> 204
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 2518753CD1
<400> 17
Met Ala Lys Val Gln Val Asn Asn Val Val Leu Asp Asn Pro
                                      10
Ser Pro Phe Tyr Asn Pro Phe Gln Phe Glu Ile Thr Phe Glu Cys
                 20
                                                           30
Ile Glu Asp Leu Ser Glu Asp Leu Glu Trp Lys Ile Ile Tyr Val
                 35
                                      40
Gly Ser Ala Glu Ser Glu Glu Tyr Asp Gln Val Leu Asp Ser Val
                 50
                                      55
Leu Val Gly Pro Val Pro Ala Gly Arg His Met Phe Val Phe Gln
                 65
                                      70
Ala Asp Ala Pro Asn Pro Gly Leu Ile Pro Asp Ala Asp Ala Val
                 80
                                      85
Gly Val Thr Val Val Leu Ile Thr Cys Thr Tyr Arg Gly Gln Glu
                 95
                                     100
Phe Ile Arg Val Gly Tyr Tyr Val Asn Asn Glu Tyr Thr Glu Thr
                110
                                     115
                                                         120
Glu Leu Arg Glu Asn Pro Pro Val Lys Pro Asp Phe Ser Lys Leu
                125
                                     130
Gln Arg Asn Ile Leu Ala Ser Asn Pro Arg Val Thr Arg Phe His
                140
                                     145
Ile Asn Trp Glu Asp Asn Thr Glu Lys Leu Glu Asp Ala Glu Ser
                155
                                     160
Ser Asn Pro Asn Leu Gln Ser Leu Leu Ser Thr Asp Ala Leu Pro
                170
                                     175
Ser Ala Ser Lys Gly Trp Ser Thr Ser Glu Asn Ser Leu Asn Val
                185
```

Met Leu Glu Ser His Met Asp Cys Met

<210> 18 <211> 713 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte clone 2709055CD1 <400> 18 Met Tyr Leu Leu Ile Gln Met Cys Tyr His Leu Ala Leu Pro Trp Tyr Ser Lys Tyr Phe Pro Tyr Leu Ala Leu Ile His Thr Ile Ile Leu Met Ala Ser Ser Asn Phe Trp Phe Lys Tyr Pro Lys Thr Cys Ser Lys Val Glu His Ser Val Ser Ile Leu Gly Lys Cys Phe Glu Ser Pro Trp Thr Thr Lys Ala Leu Ser Glu Thr Ala Cys Glu Asp Ser Glu Glu Asn Lys Gln Arg Ile Thr Gly Ala Gln Thr Leu Pro Lys His Val Ser Thr Ser Ser Asp Glu Gly Ser Pro Ser Ala Ser Thr Pro Met Ile Asn Lys Thr Gly Phe Lys Phe Ser Ala Glu Lys Pro Val Ile Glu Val Pro Ser Met Thr Ile Leu Asp Lys Lys Asp Gly Glu Gln Ala Lys Ala Leu Phe Glu Lys Val Arg Lys Phe Arg Ala His Val Glu Asp Ser Asp Leu Ile Tyr Lys Leu Tyr Val Val Gln Thr Val Ile Lys Thr Ala Lys Phe Ile Phe Ile Leu Cys Tyr Thr Ala Asn Phe Val Asn Ala Ile Ser Phe Glu His Val Cys Lys Pro Lys Val Glu His Leu Ile Gly Tyr Glu Val Phe Glu Cys Thr His Asn Met Ala Tyr Met Leu Lys Lys Leu Leu Ile Ser Tyr Ile Ser Ile Ile Cys Val Tyr Gly Phe Ile Cys Leu Tyr Thr Leu Phe Trp Leu Phe Arg Ile Pro Leu Lys Glu Tyr Ser Phe Glu Lys Val Arg Glu Glu Ser Ser Phe Ser Asp Ile Pro Asp Val Lys Asn Asp Phe Ala Phe Leu Leu His Met Val Asp Gln Tyr Asp Gln Leu Tyr Ser Lys Arg Phe Gly Val Phe Leu Ser Glu Val Ser Glu Asn Lys Leu Arg Glu Ile Ser Leu Asn His Glu Trp Thr Phe Glu Lys Leu Arg Gln His Ile Ser Arg Asn Ala Gln Asp Lys Gln Glu Leu His Leu Phe Met Leu Ser Gly Val Pro Asp Ala Val Phe Asp Leu Thr Asp Leu Asp Val Leu Lys Leu Glu Leu Ile Pro Glu Ala Lys Ile Pro Ala Lys Ile Ser Gln Met Thr Asn Leu Gln Glu Leu His Leu Cys His Cys Pro Ala Lys Val Glu Gln Thr Ala Phe Ser Phe Leu

```
380
                                      385
                                                           390
Arg Asp His Leu Arg Cys Leu His Val Lys Phe Thr Asp Val Ala
                 395
                                      400
Glu Ile Pro Ala Trp Val Tyr Leu Leu Lys Asn Leu Arg Glu Leu
                 410
                                      415
Tyr Leu Ile Gly Asn Leu Asn Ser Glu Asn Asn Lys Met Ile Gly
                 425
                                      430
                                                           435
Leu Glu Ser Leu Arg Glu Leu Arg His Leu Lys Ile Leu His Val
                 440
                                      445
Lys Ser Asn Leu Thr Lys Val Pro Ser Asn Ile Thr Asp Val Ala
                 455
                                      460
Pro His Leu Thr Lys Leu Val Ile His Asn Asp Gly Thr Lys Leu
                 470
                                     475
Leu Val Leu Asn Ser Leu Lys Lys Met Met Asn Val Ala Glu Leu
                 485
                                     490
Glu Leu Gln Asn Cys Glu Leu Glu Arg Ile Pro His Ala Ile Phe
                500
                                     505
                                                          510
Ser Leu Ser Asn Leu Gln Glu Leu Asp Leu Lys Ser Asn Asn Ile
                515
                                     520
                                                          525
Arg Thr Ile Glu Glu Ile Ile Ser Phe Gln His Leu Lys Arg Leu
                530
                                     535
Thr Cys Leu Lys Leu Trp His Asn Lys Ile Val Thr Ile Pro Pro
                545
                                     550
Ser Ile Thr His Val Lys Asn Leu Glu Ser Leu Tyr Phe Ser Asn
                560
                                     565
                                                          570
Asn Lys Leu Glu Ser Leu Pro Val Ala Val Phe Ser Leu Gln Lys
                575
                                     580
Leu Arg Cys Leu Asp Val Ser Tyr Asn Asn Ile Ser Met Ile Pro
                590
                                     595
Ile Glu Ile Gly Leu Leu Gln Asn Leu Gln His Leu His Ile Thr
                605
                                     610
Gly Asn Lys Val Asp Ile Leu Pro Lys Gln Leu Phe Lys Cys Ile
                620
                                     625
                                                          630
Lys Leu Arg Thr Leu Asn Leu Gly Gln Asn Cys Ile Thr Ser Leu
                635
                                     640
Pro Glu Lys Val Gly Gln Leu Ser Gln Leu Thr Gln Leu Glu Leu
                650
                                     655
                                                          660
Lys Gly Asn Cys Leu Asp Arg Leu Pro Ala Gln Leu Gly Gln Cys
                665
                                     670
                                                          675
Arg Met Leu Lys Lys Ser Gly Leu Val Val Glu Asp His Leu Phe
                680
                                     685
Asp Thr Leu Pro Leu Glu Val Lys Glu Ala Leu Asn Gln Asp Ile
                695
                                     700
                                                          705
Asn Ile Pro Phe Ala Asn Gly Ile
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<210> 19

<211> 360

<212> PRT

<213> Homo sapiens

<220>

<221> misc\_feature

<223> Incyte clone 2724537CD1

<400> 19

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50
                                     55
                                                          60
Lys Ala Ala Glu His Lys Ala Lys Ser Leu Gly Glu Lys Ser Pro
                 65
                                     70
Ala Ala Ser Gly Ala Arg Arg Pro Glu Ala Ala Lys Glu Glu Ala
                                     85
Ala Trp Ala Ser Ser Ser Ala Gly Asn Pro Ala Asp Gly Leu Ala
                 95
                                    100
Thr Glu Pro Glu Ser Val Phe Ala Leu Asp Val Leu Arg Gln Arg
                110
                                    115
                                                         120
Leu His Glu Lys Ile Gln Glu Ala Arg Gly Gln Gly Ser Ala Lys
                125
                                    130
                                                         135
Glu Leu Ser Pro Ala Ala Leu Glu Lys Arg Arg Arg Lys Gln
                140
                                    145
Glu Arg Asp Arg Lys Lys Arg Lys Glu Leu Arg Ala Lys
                155
                                    160
Glu Lys Ala Arg Lys Ala Glu Glu Ala Thr Glu Ala Gln Glu Val
                170
                                    175
Val Glu Ala Thr Pro Glu Gly Ala Cys Thr Glu Pro Arg Glu Pro
                185
                                    190
                                                         195
Pro Gly Leu Ile Phe Asn Lys Val Glu Val Ser Glu Asp Glu Pro
                200
                                    205
                                                         210
Ala Ser Lys Ala Gln Arg Arg Lys Glu Lys Arg Gln Arg Val Lys
                215
                                    220
                                                         225
Gly Asn Leu Thr Pro Leu Thr Gly Arg Asn Tyr Arg Gln Leu Leu
                230
                                    235
                                                         240
Glu Arg Leu Gln Ala Arg Gln Ser Arg Leu Asp Glu Leu Arg Gly
                245
                                    250
                                                         255
Gln Asp Glu Gly Lys Ala Gln Glu Leu Glu Ala Lys Met Lys Trp
                260
                                    265
                                                         270
Thr Asn Leu Leu Tyr Lys Ala Glu Gly Val Lys Ile Arg Asp Asp
                275
                                    280
                                                         285
Glu Arg Leu Leu Gln Glu Ala Leu Lys Arg Lys Glu Lys Arg Arg
                290
                                    295
                                                         300
Ala Gln Arg Gln Arg Trp Glu Lys Arg Thr Ala Gly Val Val
                305
                                    310
                                                         315
Glu Lys Met Gln Gln Arg Gln Asp Arg Arg Gln Asn Leu Arg
                320
                                    325
                                                         330
Arg Lys Lys Ala Ala Arg Ala Glu Arg Arg Leu Leu Arg Ala Arg
                335
                                    340
Lys Lys Gly Arg Ile Leu Pro Gln Asp Leu Glu Arg Ala Gly Leu
                350
```

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<210> 20
<211> 196
<212> PRT
<213> Homo sapiens
<220>
<221> misc feature
<223> Incyte clone 025818CD1
<400> 20
Met Pro Ala Asp Ile Met Glu Lys Asn Ser Ser Ser Pro Val Ala
                                     10
Ala Thr Pro Ala Ser Val Asn Thr Thr Pro Asp Lys Pro Lys Thr
                                                          30
Ala Ser Glu His Arg Lys Ser Ser Lys Pro Ile Met Glu Lys Arg
                 35
Arg Arg Ala Arg Ile Asn Glu Ser Leu Ser Gln Leu Lys Thr Leu
                 50
                                      55
Ile Leu Asp Ala Leu Lys Lys Asp Ser Ser Arg His Ser Lys Leu
```

```
65
                                      70
Glu Lys Ala Asp Ile Leu Glu Met Thr Val Lys His Leu Arg Asn
                 80
                                      85
Leu Gln Arg Ala Gln Met Thr Ala Ala Leu Ser Thr Asp Pro Ser
                                     100
                                                          105
Val Leu Gly Lys Tyr Arg Ala Gly Phe Ser Glu Cys Met Asn Glu
                110
                                     115
                                                          120
Val Thr Arg Phe Leu Ser Ser Pro Ser Thr Pro Ala Thr Ala Ala
                125
                                     130
                                                          135
Pro Pro Trp Ala Pro Thr Gln Cys His Leu Pro Ala Ala Pro Arg
                140
                                     145
                                                          150
Leu Arg Arg Thr Pro Cys Gly Gly Arg Gly Gly Thr Glu Gly Ala
                155
                                     160
Gln Ala Thr Pro Pro Pro Lys Leu Pro Asn Pro Pro Leu Phe Pro
                170
                                     175
Pro Asp Ser Lys Gln Glu Leu Glu Tyr Trp Glu Arg Arg Gly Leu
                185
                                     190
Phe
```

<210> 21

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<211> 540
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 438283CD1
<400> 21
Met Leu Arg Glu Glu Ala Thr Lys Lys Ser Lys Glu Lys Glu Pro
                                      10
Gly Met Ala Leu Pro Gln Gly Arg Leu Ala Phe Arg Asp Val Ala
                 20
                                      25
Ile Glu Phe Ser Leu Glu Glu Trp Lys Cys Leu Asn Pro Ala Gln
                 35
                                      40
                                                           45
Arg Ala Leu Tyr Arg Ala Val Met Leu Glu Asn Tyr Arg Asn Leu
                 50
                                      55
Glu Phe Val Asp Ser Ser Leu Lys Ser Met Met Glu Phe Ser Ser
                 65
                                      70
                                                           75
Thr Arg His Ser Asn Thr Gly Glu Val Ile His Thr Gly Thr Leu
                 80
                                      8.5
Gln Arg His Lys Ser His His Ile Gly Asp Phe Cys Phe Pro Glu
                 95
                                     100
Met Lys Lys Asp Ile His His Phe Glu Phe Gln Trp Gln Glu Val
                110
                                     115
Glu Arg Asn Gly His Glu Ala Pro Met Thr Lys Ile Lys Lys Leu
                125
                                     130
                                                          135
Thr Gly Ser Thr Asp Arg Ser Asp His Arg His Ala Gly Asn Lys
                140
                                     145
Pro Ile Lys Asp Gln Leu Gly Leu Ser Phe His Ser His Leu Pro
                155
                                     160
                                                          165
Glu Leu His Met Phe Gln Thr Lys Gly Lys Ile Ser Asn Gln Leu
                170
                                     175
Asp Lys Ser Ile Ser Gly Ala Ser Ser Ala Ser Glu Ser Gln Arg
                185
                                     190
Ile Ser Cys Arg Leu Lys Thr His Ile Ser Asn Lys Tyr Gly Lys
                200
                                     205
                                                          210
Asn Phe Leu His Ser Ser Phe Thr Gln Ile Gln Glu Ile Cys Met
                215
                                     220
Arg Glu Lys Pro Cys Gln Ser Asn Glu Cys Gly Lys Ala Phe Asn
                230
                                     235
```

```
Tyr Ser Ser Leu Leu Arg Arg His His Ile Thr His Ser Arg Glu
                 245
                                      250
                                                          255
Arg Glu Tyr Lys Cys Asp Val Cys Gly Lys Ile Phe Asn Gln Lys
                 260
                                      265
Gln Tyr Ile Val Tyr His His Arg Cys His Thr Gly Glu Lys Thr
                 275
                                      280
                                                          285
Tyr Lys Cys Asn Glu Cys Gly Lys Thr Phe Thr Gln Met Ser Ser
                 290
                                      295
                                                          300
Leu Val Cys His Arg Arg Leu His Thr Gly Glu Lys Pro Tyr Lys
                 305
                                     310
                                                          315
Cys Asn Glu Cys Gly Lys Thr Phe Ser Glu Lys Ser Ser Leu Arg
                 320
                                     325
                                                          330
Cys His Arg Arg Leu His Thr Gly Glu Lys Pro Tyr Lys Cys Asn
                 335
                                     340
Glu Cys Gly Lys Thr Phe Gly Arg Asn Ser Ala Leu Val Ile His
                 350
                                     355
                                                          360
Lys Ala Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Asn Glu Cys
                 365
                                     370
                                                          375
Gly Lys Thr Phe Ser Gln Lys Ser Ser Leu Gln Cys His His Ile
                380
                                     385
                                                          390
Leu His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys Asp Asn
                395
                                     400
                                                          405
Val Tyr Ile Arg Arg Ser His Leu Glu Arg His Arg Lys Ile His
                410
                                     415
Thr Gly Glu Gly Ser Tyr Lys Cys Lys Val Cys Asp Lys Ala Phe
                425
                                     430
                                                          435
Arg Ser Asp Ser Cys Leu Ala Asn His Thr Arg Val His Thr Gly
                440
                                     445
                                                          450
Glu Lys Pro Tyr Lys Cys Asn Lys Cys Ala Lys Val Phe Asn Gln
                455
                                     460
                                                          465
Lys Gly Ile Leu Ala Gln His Gln Arg Val His Thr Gly Glu Lys
                470
                                     475
Pro Tyr Lys Cys Asn Glu Cys Gly Lys Val Phe Asn Gln Lys Ala
                485
                                     490
                                                          495
Ser Leu Ala Lys His Gln Arg Val His Thr Ala Glu Lys Pro Tyr
                500
                                     505
Lys Cys Asn Glu Cys Gly Lys Ala Phe Thr Gly Gln Ser Thr Leu
                515
                                     520
                                                          525
Ile His His Gln Ala Ile His Gly Cys Arg Glu Thr Leu Gln Met
                530
                                     535
```

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<210> 22
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## <400> 22

Met Leu Glu Asn Tyr Lys Asn Leu Ala Thr Val Gly Tyr Gln Leu Phe Lys Pro Ser Leu Ile Ser Trp Leu Glu Glu Glu Ser Arg Thr Val Gln Arg Gly Asp Phe Gln Ala Ser Glu Trp Lys Val Gln 4Ω Leu Lys Thr Lys Glu Leu Ala Leu Gln Gln Asp Val Leu Gly Glu Pro Thr Ser Ser Gly Ile Gln Met Ile Gly Ser His Asn Gly Gly Glu Val Ser Asp Val Lys Gln Cys Gly Asp Val Ser Ser Glu His

<sup>&</sup>lt;211> 549

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

<sup>&</sup>lt;220>

<sup>&</sup>lt;221> misc\_feature

<sup>&</sup>lt;223> Incyte clone 619699CD1

				9:	)				1 0 0	1				Thr
Ph∈	Glu	Cys د	з Ту	r Leu 110	ı Tyr	Gly	/ Val	Asp	Phe 115	. Le	ı Thı	Leu	His	Lys 120
				125	)				Val	Phe				Gly
Lys	Ala	Phe	e Se	r Let 140	Asn	Pro	Asp	Val	. Val	Cys	Glr	n Arg	Thr	Cys 150
Thr	Gl	/ Glu	ı Lys	s Ala 155	Phe	Asp	Cys	Ser	Asp 160	Ser	: Gly	/ Lys	Ser	Phe: 165
Ile	Asr	n His	Se		Leu	Gln	Gly	His	Leu 175	Arg	Thr	His	Asn	Gly
Glu	Ser	Leu	His		Trp	Lys	Glu	Cys	Gly 190	Arg	Gly	Phe	Ile	180 His
Ser	Thr	Asp	Lei		Val	Arg	Ile	Gln	Thr 205	His	Arg	Ser	Glu	195 Lys
Pro	Tyr	Lys	Cys		Glu	Cys	Gly	Lys	Gly 220	Phe	Arg	Tyr	Ser	210 Ala
Tyr	Leu	Asn	Ile		Met	Gly	Thr	His	Thr 235	Gly	Asp	Asn	Pro	225 Tyr
Glu	Cys	Lys	Glu		Gly	Lys	Ala	Phe	Thr 250	Arg	Ser	Cys	Gln	240 Leu 255
Thr	Gln	His	Arg		Thr	His	Thr	Gly	Glu 265	Lys	Pro	Tyr	Lys	Cys
Lys	Asp	Cys	Gl <sub>y</sub>	Arg 275	Ala	Phe	Thr	Val	Ser 280	Ser	Cys	Leu	Ser	270 Gln 285
His	Met	Lys	Ile	His 290	Val	Gly	Glu	Lys	Pro 295	Tyr	Glu	Cys	Lys	Glu 300
Cys	Gly	Ile	Ala	Phe 305	Thr	Arg	Ser	Ser	Gln 310	Leu	Thr	Glu	His	Leu 315
Lys	Thr	His	Thr	Ala 320	Lys	Asp	Pro	Phe	Glu 325	Cys	Lys	Val	Cys	Gly 330
Lys	Ser	Phe	Arg	Asn 335	Ser	Ser	Cys	Leu	Ser 340	Asp	His	Phe	Arg	Ile 345
His	Thr	Gly	Ile	Lys 350	Pro	Tyr	Lys	Cys	Lys 355	Asp	Cys	Gly	Lys	Ala 360
Phe	Thr	Gln	Asn	Ser 365	Asp	Leu	Thr	Lys	His 370	Ala	Arg	Thr	His	Ser 375
				Tyr 380					Cys 385					Ala
				Leu 395					Arg					Glu 405
				Cys 410					Lys					Ser
Ser	Asn	Leu	Ser	Gly 425	His	Leu	Arg	Ile	His 430	Thr	Gly	Glu	Lys	Pro 435
				Glu 440					Phe					Ser
Leu	Asn	Asn	His	Met 455	Arg	Thr	His	Ser	Ala 460	Lys	Lys	Pro	Phe	Thr 465
Cys	Met	Glu	Cys	Gly 470	Lys	Ala	Phe	Lys	Phe 475	Pro	Thr	Cys	Val	Asn 480
				Ile 485					Lys 490					Lys
Gln	Cys	Gly	Lys	Ser 500	Phe	Ser	Tyr	Ser	Asn 505	Ser	Phe	Gln	Leu	His 510
Glu	Arg	Thr	His	Thr 515	Gly	Glu	Lys	Pro	Tyr 520	Glu	Cys	Lys	Glu	Cys 525
Gly	Lys	Ala	Phe	Ser 530	Ser	Ser	Ser	Ser	Phe 535	Arg	Asn	His	Glu	Arg
Arg	His	Ala	Asp	Glu 545	Arg	Leu	Ser	Ala						540

<210> 23

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<211> 361
 <212> PRT
 <213> Homo sapiens
<220>
 <221> misc feature
<223> Incyte clone 693452CD1
<400> 23
Met Ala Asp Phe Lys Val Leu Ser Ser Gln Asp Ile Lys Trp Ala
                                      10
Leu His Glu Leu Lys Gly His Tyr Ala Ile Thr Arg Lys Ala Leu
                  20
                                       25
Ser Asp Ala Ile Lys Lys Trp Gln Glu Leu Ser Pro Glu Thr Ser
                  35
                                       40
Gly Lys Arg Lys Arg Lys Gln Met Asn Gln Tyr Ser Tyr Ile
                  50
                                      55
Asp Phe Lys Phe Glu Gln Gly Asp Ile Lys Ile Glu Lys Arg Met
                 65
Phe Phe Leu Glu Asn Lys Arg Arg His Cys Arg Ser Tyr Asp Arg
                 80
                                      85
Arg Ala Leu Leu Pro Ala Val Gln Gln Glu Gln Glu Phe Tyr Glu
                 95
                                     100
                                                          105
Gln Lys Ile Lys Glu Met Ala Glu His Glu Asp Phe Leu Leu Ala
                110
                                     115
                                                          120
Leu Gln Met Asn Glu Glu Gln Tyr Gln Lys Asp Gly Gln Leu Ile
                125
                                     130
                                                          135
Glu Cys Arg Cys Cys Tyr Gly Glu Phe Pro Phe Glu Glu Leu Thr
                140
                                     145
Gln Cys Ala Asp Ala His Leu Phe Cys Lys Glu Cys Leu Ile Arg
                155
                                     160
                                                          165
Tyr Ala Gln Glu Ala Val Phe Gly Ser Gly Lys Leu Glu Leu Ser
                170
                                     175
                                                          180
Cys Met Glu Gly Ser Cys Thr Cys Ser Phe Pro Thr Ser Glu Leu
                185
                                     190
                                                          195
Glu Lys Val Leu Pro Gln Thr Ile Leu Tyr Lys Tyr Tyr Glu Arg
                200
                                     205
Lys Ala Glu Glu Val Ala Ala Ala Tyr Ala Asp Glu Leu Val
                215
                                     220
                                                          225
Arg Cys Pro Ser Cys Ser Phe Pro Ala Leu Leu Asp Ser Asp Val
                230
                                     235
                                                          240
Lys Arg Phe Ser Cys Pro Asn Pro His Cys Arg Lys Glu Thr Cys
                245
                                     250
                                                          255
Arg Lys Cys Gln Gly Leu Trp Lys Glu His Asn Gly Leu Thr Cys
                260
                                     265
Glu Glu Leu Ala Glu Lys Asp Asp Ile Lys Tyr Arg Thr Ser Ile
                275
                                     280
Glu Glu Lys Met Thr Ala Ala Arg Ile Arg Lys Cys His Lys Cys
                290
                                     295
                                                          300
Gly Thr Gly Leu Ile Lys Ser Glu Gly Cys Asn Arg Met Ser Cys
                305
                                     310
Arg Cys Gly Ala Gln Met Cys Tyr Leu Cys Arg Val Ser Ile Asn
                320
                                     325
                                                          330
Gly Tyr Asp His Xaa Cys Gln Gln Ser Arg Leu Thr Gly Ala Pro
                                     340
Phe Gln Gly Val Phe Lys Met Leu Ser Met Asp Arg Leu Gln Cys
                350
                                     355
                                                         360
Lys
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<210> 24
<211> 241
<212> PRT
<213> Homo sapiens
<220>
<221> misc feature
<223> Incyte clone 839651CD1
<400> 24
Met Trp Pro Ser Leu Glu Ala Leu Cys Ser Leu Phe Ala Ala Arg
                                      10
Ser Thr Gly Ser Gln Ala Gln Ser Ala Pro Thr Pro Ala Trp Asp
                  20
                                      25
                                                           30
Glu Asp Thr Ala Gln Ile Gly Pro Lys Arg Ile Arg Lys Ala Ala
                  35
                                      40
Lys Arg Glu Leu Met Pro Cys Asp Phe Pro Gly Cys Gly Arg Ile
                 50
                                      55
Phe Ser Asn Arg Gln Tyr Leu Asn His His Lys Lys Tyr Gln His
                 65
                                      70
Ile His Gln Lys Ser Phe Ser Cys Pro Glu Pro Ala Cys Gly Lys
                 80
                                      85
                                                           90
Ser Phe Asn Phe Lys Lys His Leu Lys Glu His Met Lys Leu His
                 95
                                     100
                                                          105
Ser Asp Thr Arg Asp Tyr Ile Cys Glu Phe Cys Ala Arg Ser Phe
                110
                                     115
                                                          120
Arg Thr Ser Ser Asn Leu Val Ile His Arg Arg Ile His Thr Gly
                125
                                     130
Glu Lys Pro Leu Gln Cys Glu Ile Cys Gly Phe Thr Cys Arg Gln
                140
                                     145
Lys Ala Ser Leu Asn Trp His Gln Arg Lys His Ala Glu Thr Val
                155
                                     160
                                                          165
Ala Ala Leu Arg Phe Pro Cys Glu Phe Cys Gly Lys Arg Phe Glu
                170
                                     175
                                                          180
Lys Pro Asp Ser Val Ala Ala His Arg Ser Lys Ser His Pro Ala
                185
                                     190
Leu Leu Leu Ala Pro Gln Glu Ser Pro Ser Gly Pro Leu Glu Pro
                200
                                     205
Cys Pro Ser Ile Ser Ala Pro Gly Pro Leu Gly Ser Ser Glu Gly
                215
                                     220
                                                          225
Ser Arg Pro Ser Ala Ser Pro Gln Ala Pro Thr Leu Leu Pro Gln
                230
                                     235
                                                          240
Gln
```

<210> 25 <211> 576

				50					55					60
Lys	Glu	Ile	Asn	Gly 65	Ile	His	Asp	Glu	Ser 70	Asn	Ala	Phe	Glu	Ser 75
Ī			Glu	80			Leu		85					Ser 90
Gln	Phe	Phe	Glu	Gln 95	Gly	Ser	Ser	Asp	Ser 100	Val	Val	Pro	Asp	Leu 105
Pro	Val	Pro	Thr	Ile 110	Ser	Ala	Pro	Ser	Arg 115	Trp	Val	Trp	Asp	
Glu	Glu	Glu	Arg	Lys 125	Arg	Gln	Glu	Arg	Trp	Gln	Lys	Glu	Gln	Asp
Arg	Leu	Leu	Gln	Glu 140	Lys	Tyr	Gln	Arg	Glu 145	Gln	Glu	Lys	Leu	Arg 150
			Gln	155					160					Ser 165
			Asp	170					175					Met 180
			Thr	185					190					195
			Gly	200					205					210
			Glu	215					220					225
			Lys	230					235				_	240
			Glu	245					250					255
			Glu	260					265					270
_	_		Ala	275			Arg		280					285
			Arg	290					295					300
			Ser	305					310					315
			Glu	320					325				_	330
			Leu	335					340					345
	Ser		Lys	350					355					360
			Leu	365					370					375
			Trp	380					385				=	390
			Leu	395			-		400					405
			Ser	410					415					420
			Pro Pro	425					430					435
			Asn	440					445				-	450
			Thr	455					460					465
			Ser	470					475					Pro 480
			Arg	485					490					495
			Leu	500					505					510
			Tyr	515					520					525
y		J, J	- Y -		⊿.eu	.113	CAP	rne	гÃ2	Cys	vdl	Ата	cys	ern

```
530
                                     535
Cys Asp Leu Gly Gly Ser Ser Ser Gly Ala Glu Val Arg Ile Arg
                 545
                                     550
 Asn His Gln Leu Tyr Cys Asn Asp Cys Tyr Leu Arg Phe Lys Ser
                 560
                                     565
Gly Arg Pro Thr Ala Met
                 575
 <210> 26
 <211> 408
 <212> PRT
<213> Homo sapiens
<220>
<221> misc feature
<223> Incyte clone 1425691CD1
<400> 26
Met Pro Gly His Leu Gln Glu Gly Phe Gly Cys Val Val Thr Asn
                                      10
Arg Phe Asp Gln Leu Phe Asp Asp Glu Ser Asp Pro Phe Glu Val
                                      25
                                                           30
Leu Lys Ala Ala Glu Asn Lys Lys Glu Ala Gly Gly Gly
Val Gly Gly Pro Gly Ala Lys Ser Ala Ala Gln Ala Ala Gln
                 50
                                      55
Thr Asn Ser Asn Ala Ala Gly Lys Gln Leu Arg Lys Glu Ser Gln
                 65
                                      70
Lys Asp Arg Lys Asn Pro Leu Pro Pro Ser Val Gly Val Val Asp
                 80
                                      85
Lys Lys Glu Glu Thr Gln Pro Pro Val Ala Leu Lys Lys Glu Gly
                 95
                                     100
                                                         105
Ile Arg Arg Val Gly Arg Arg Pro Asp Gln Gln Leu Gln Gly Glu
                110
                                    115
                                                         120
Gly Lys Ile Ile Asp Arg Arg Pro Glu Arg Arg Pro Pro Arg Glu
                125
                                     130
Arg Arg Phe Glu Lys Pro Leu Glu Glu Lys Gly Glu Gly Glu
                140
                                     145
Phe Ser Val Asp Arg Pro Ile Ile Asp Arg Pro Ile Arg Gly Arg
                155
                                     160
                                                         165
Gly Gly Leu Gly Arg Gly Arg Gly Arg Gly Arg Gly Met Gly
                170
                                     175
                                                         180
Arg Gly Asp Gly Phe Asp Ser Arg Gly Lys Arg Glu Phe Asp Arg
                185
                                    190
His Ser Gly Ser Asp Arg Ser Ser Phe Ser His Tyr Ser Gly Leu
                200
                                    205
Lys His Glu Asp Lys Arg Gly Gly Ser Gly Ser His Asn Trp Gly
                215
                                     220
Thr Val Lys Asp Glu Leu Thr Glu Ser Pro Lys Tyr Ile Gln Lys
                230
                                    235
Gln Ile Ser Tyr Asn Tyr Ser Asp Leu Asp Gln Ser Asn Val Thr
                245
                                    250
                                                         255
Glu Glu Thr Pro Glu Gly Glu Glu His His Pro Val Ala Asp Thr
                260
                                    265
Glu Asn Lys Glu Asn Glu Val Glu Glu Val Lys Glu Glu Gly Pro
                275
                                     280
                                                         285
Lys Glu Met Thr Leu Asp Glu Trp Lys Ala Ile Gln Asn Lys Asp
                290
                                     295
                                                         300
Arg Ala Lys Val Glu Phe Asn Ile Arg Lys Pro Asn Glu Gly Ala
                305
                                    310
Asp Gly Gln Trp Lys Lys Gly Phe Val Leu His Lys Ser Lys Ser
                320
                                    325
Glu Glu Ala His Ala Glu Asp Ser Val Met Asp His His Phe Arg
```

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335
                                     340
                                                          345
 Lys Pro Ala Asn Asp Ile Thr Ser Gln Leu Glu Ile Asn Phe Gly
                 350
                                      355
                                                          360
 Asp Leu Gly Arg Pro Gly Arg Gly Gly Arg Gly Arg Gly Gly
                 365
                                      370
                                                          375
 Arg Gly Arg Gly Gly Arg Pro Asn Arg Gly Ser Arg Thr Asp Lys
                 380
                                     385
                                                          390
 Ser Ser Ala Ser Ala Pro Asp Val Asp Asp Pro Glu Ala Phe Pro
                 395
                                     400
Ala Leu Ala
<210> 27
<211> 810
<212> PRT
<213> Homo sapiens
<220>
<221> misc feature
<223> Incyte clone 1484257CD1
<400> 27
Met Asp Phe Pro Gln His Ser Gln His Val Leu Glu Gln Leu Asn
                                      10
Gln Gln Arg Gln Leu Gly Leu Leu Cys Asp Cys Thr Phe Val Val
                  20
                                                           30
Asp Gly Val His Phe Lys Ala His Lys Ala Val Leu Ala Ala Cys
                  35
                                      40
Ser Glu Tyr Phe Lys Met Leu Phe Val Asp Gln Lys Asp Val Val
                  50
                                      55
His Leu Asp Ile Ser Asn Ala Ala Gly Leu Gly Gln Val Leu Giu
                  65
                                      70
Phe Met Tyr Thr Ala Lys Leu Ser Leu Ser Pro Glu Asn Val Asp
                 80
                                      85
Asp Val Leu Ala Val Ala Thr Phe Leu Gln Met Gln Asp Ile Ile
                  95
                                     100
                                                          105
Thr Ala Cys His Ala Leu Lys Ser Leu Ala Glu Pro Ala Thr Ser
                110
                                     115
                                                         120
Pro Gly Gly Asn Ala Glu Ala Leu Ala Gln Lys Val Cys Pro Val
                125
                                     130
Pro Ser Pro Gly Gly Asp Lys Arg Ala Lys Glu Glu Lys Val Ala
                140
                                     145
                                                         150
Thr Ser Thr Leu Ser Arg Leu Glu Gln Ala Gly Arg Ser Thr Pro
                155
                                     160
Ile Gly Pro Ser Arg Asp Leu Lys Glu Glu Arg Gly Gln Ala
                170
                                     175
                                                         180
Gln Ser Ala Ala Ser Gly Ala Glu Gln Thr Glu Lys Ala Asp Ala
                185
                                     190
Pro Arg Glu Pro Pro Pro Val Glu Leu Lys Pro Asp Pro Thr Ser
                200
                                     205
Gly Met Ala Ala Ala Glu Ala Glu Ala Ala Leu Ser Glu Ser Ser
                215
                                     220
                                                         225
Glu Gln Glu Met Glu Val Glu Pro Ala Arg Lys Gly Glu Glu
                230
                                     235
                                                         240
Gln Lys Glu Gln Glu Glu Glu Glu Glu Gly Ala Gly Pro Ala
                245
                                     250
Glu Val Lys Glu Glu Gly Ser Gln Leu Glu Asn Gly Glu Ala Pro
                260
                                     265
Glu Glu Asn Glu Asn Glu Glu Ser Ala Gly Thr Asp Ser Gly Gln
                275
                                     280
                                                         285
Glu Leu Gly Ser Glu Ala Arg Gly Leu Arg Ser Gly Thr Tyr Gly
                290
                                     295
                                                         300
Asp Arg Thr Glu Ser Lys Ala Tyr Gly Ser Val Ile His Lys Cys
                305
                                     310
                                                         315
```

Glu	Asp	Cys	Gly	Lys 320	Glu	Phe	Thr	His	Thr 325	Gly	Asn	Phe	. Lys	Arg 330
His	Ile	Arg	ılle	His 335		Gly	Glu	Lys		Phe	Ser	Cys	Arg	Glu 345
				350					355				His	Glu 360
				365					370				Cys	Gly 375
			Arg	380					385				Lys	390
				395					400				Lys	405
				410					415				His	420
				425					430				Phe	435
				440					445				Thr	Asp
				455					460				Gln	465
				470					475				Gly	480
				485					490				Gly	495
				500					505				Tyr	Val
				515					520				Leu	Gln
				530					535				Cys	540
				545					550	Ser	Leu	Ile	Ala	His 555
				560			Lys		565		_		Arg	Cys
				575					580				Ile	Arg 585
				590			His		595	Ser	Val	Cys	Ser	Lys 600
				605			Leu		610				Ile	615
				620					625				Gly	630
				635					640				His	645
				650					655				Ser	Glu 660
				665					670				Thr	Glu 675
				680					685				Pro	690
				695					700				Ala	Glu 705
				110					715				Asn	720
				125					730				Leu	735
				740					745				Gln	750
				155					760				Tyr	Gly 765
				//0					Val 775				Gly	Glu 780
Leu	Val	Phe	Arg	Pro 785	Arg	Asp	Gly	Ala	Glu 790	Gly	Gln	Pro	Ala	Leu 795

Ala Glu Thr Ser Pro Thr Ala Pro Glu Cys Pro Pro Pro Ala Glu 800 805 805

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<210> 28
<211> 324
<212> PRT
<213> Homo sapiens
<220>
<221> misc feature
<223> Incyte clone 1732368CD1
<400> 28
Met Asp Trp Ser Glu Val Lys Glu Glu Lys Asp Asn Leu Glu Ile
                                      10
Lys Gln Glu Glu Lys Phe Val Gly Gln Cys Ile Lys Glu Glu Leu
                 20
Met His Gly Glu Cys Val Lys Glu Glu Lys Asp Phe Leu Lys Lys
                                      40
                                                           45
Glu Ile Val Asp Asp Thr Lys Val Lys Glu Glu Pro Pro Ile Asn
                 50
His Pro Val Gly Cys Lys Arg Lys Leu Ala Met Ser Arg Cys Glu
                 65
                                      70
Thr Cys Gly Thr Glu Glu Ala Lys Tyr Arg Cys Pro Arg Cys Met
                 80
                                      85
Arg Tyr Ser Cys Ser Leu Pro Cys Val Lys Lys His Lys Ala Glu
                 95
                                     100
Leu Thr Cys Asn Gly Val Arg Asp Lys Thr Ala Tyr Ile Ser Ile
                110
                                     115
                                                         120
Gln Gln Phe Thr Glu Met Asn Leu Leu Ser Asp Tyr Arg Phe Leu
                125
                                     130
                                                         135
Glu Asp Val Ala Arg Thr Ala Asp His Ile Ser Arg Asp Ala Phe
                140
                                     145
Leu Lys Arg Pro Ile Ser Asn Lys Tyr Met Tyr Phe Met Lys Asn
                                     160
                                                         165
Arg Ala Arg Arg Gln Gly Ile Asn Leu Lys Leu Leu Pro Asn Gly
                170
                                     175
                                                         180
Phe Thr Lys Arg Lys Glu Asn Ser Thr Phe Phe Asp Lys Lys
                185
                                     190
                                                         195
Gln Gln Phe Cys Trp His Val Lys Leu Gln Phe Pro Gln Ser Gln
                200
                                     205
Ala Glu Tyr Ile Glu Lys Arg Val Pro Asp Asp Lys Thr Ile Asn
                215
                                     220
Glu Ile Leu Lys Pro Tyr Ile Asp Pro Glu Lys Ser Asp Pro Val
                230
                                     235
Ile Arg Gln Arg Leu Lys Ala Tyr Ile Arg Ser Gln Thr Gly Val
                245
                                    250
                                                         255
Gln Ile Leu Met Lys Ile Glu Tyr Met Gln Gln Asn Leu Val Arg
                260
                                    265
                                                         270
Tyr Tyr Glu Leu Asp Pro Tyr Lys Ser Leu Leu Asp Asn Leu Arg
                275
                                    280
                                                         285
Asn Lys Val Ile Ile Glu Tyr Pro Thr Leu His Val Val Leu Lys
                290
                                     295
                                                         300
Gly Ser Asn Asn Asp Met Lys Val Leu His Gln Val Lys Ser Glu
                305
                                    310
Ser Thr Lys Asn Val Gly Asn Glu Asn
                320
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<210> 29 <211> 292

<212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte clone 1870914CD1 <400> 29 Met Glu Glu Val Pro His Asp Cys Pro Gly Ala Asp Ser Ala Gln 10 Ala Gly Arg Gly Ala Ser Cys Gln Gly Cys Pro Asn Gln Arg Leu 20 25 Cys Ala Ser Gly Ala Gly Ala Thr Pro Asp Thr Ala Ile Glu Glu 35 40 Ile Lys Glu Lys Met Lys Thr Val Lys His Lys Ile Leu Val Leu 50 55 60 Ser Gly Lys Gly Gly Val Gly Lys Ser Thr Phe Ser Ala His Leu 65 70 Ala His Gly Leu Ala Glu Asp Glu Asn Thr Gln Ile Ala Leu Leu 85 Asp Ile Asp Ile Cys Gly Pro Ser Ile Pro Lys Ile Met Gly Leu 95 100 105 Glu Gly Glu Gln Val His Gln Ser Gly Ser Gly Trp Ser Pro Val 110 115 120 Tyr Val Glu Asp Asn Leu Gly Val Met Ser Val Gly Phe Leu Leu 125 130 Ser Ser Pro Asp Asp Ala Val Ile Trp Arg Gly Pro Lys Lys Asn 140 145 Gly Met Ile Lys Gln Phe Leu Arg Asp Val Asp Trp Gly Glu Val 155 160 165 Asp Tyr Leu Ile Val Asp Thr Pro Pro Gly Thr Ser Asp Glu His 170 175 180 Leu Ser Val Val Arg His Leu Ala Thr Ala His Ile Asp Gly Ala 185 190 195 Val Ile Ile Thr Thr Pro Gln Glu Val Ser Leu Gln Asp Val Arg 200 205 210 Lys Glu Ile Asn Phe Cys Arg Lys Val Lys Leu Pro Ile Ile Gly 215 220 225 Val Val Glu Asn Met Ser Gly Phe Ile Cys Pro Lys Cys Lys 230 235 Glu Ser Gln Ile Phe Pro Pro Thr Thr Gly Gly Ala Glu Leu Met 245 250 Cys Gln Asp Leu Glu Val Pro Leu Leu Gly Arg Val Pro Leu Asp 260 265 270 Pro Leu Ile Gly Ile Gln Glu Phe Cys Asn Leu His Gln Ser Lys 275 280 285 Glu Glu Asn Leu Ile Ser Ser 290 <210> 30 <211> 259 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte clone 1910984CD1 <400> 30 Met Glu Cys His Leu Lys Thr His Tyr Lys Met Glu Tyr Lys Cys 10 Arg Ile Cys Gln Thr Val Lys Ala Asn Gln Leu Glu Leu Glu Thr 20

```
His Thr Arg Glu His Arg Leu Gly Asn His Tyr Lys Cys Asp Gln
Cys Gly Tyr Leu Ser Lys Thr Ala Asn Lys Leu Ile Glu His Val
                                      55
Arg Val His Thr Gly Glu Arg Pro Phe His Cys Asp Gln Cys Ser
                 65
                                      70
                                                           75
Tyr Ser Cys Thr Gly Lys Asp Asn Leu Asn Leu His Lys Lys Leu
                 80
                                      85
Lys His Ala Pro Arg Gln Thr Phe Ser Cys Glu Glu Cys Leu Phe
                 95
                                     100
                                                          105
Lys Thr Thr His Pro Phe Val Phe Ser Arg His Val Lys Lys His
                110
                                     115
Gln Ser Gly Asp Cys Pro Glu Glu Asp Lys Lys Gly Leu Cys Pro
                125
                                     130
Ala Pro Lys Glu Pro Ala Gly Pro Gly Ala Pro Leu Leu Val Val
                140
                                     145
                                                         150
Gly Ser Ser Arg Asn Leu Leu Ser Pro Leu Ser Val Met Ser Ala
                155
                                     160
Ser Gln Ala Leu Gln Thr Val Ala Leu Ser Ala Ala His Gly Ser
                170
                                     175
                                                         180
Ser Ser Glu Pro Asn Leu Ala Leu Lys Ala Leu Ala Phe Asn Gly
                185
                                     190
Ser Pro Leu Arg Phe Asp Lys Tyr Arg Asn Ser Asp Phe Ala His
                200
                                     205
                                                         210
Leu lle Pro Leu Thr Met Leu Tyr Pro Lys Asn His Leu Asp Leu
                215
                                     220
                                                         225
Thr Phe His Pro Pro Arg Pro Gln Thr Ala Pro Pro Ser Ile Pro
                230
                                     235
                                                         240
Ser Pro Lys His Ser Phe Leu Ala Tyr Leu Gly Leu Arg Glu Arg
                245
                                     250
Ala Glu Thr Val
```

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<210> 31
<211> 97
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 1943040CD1
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<400> 31 Met Glu His His Ser Ser His Gly Gly Arg Lys Arg Tyr Ala Cys 10 Gln Gly Cys Trp Lys Thr Phe His Phe Ser Leu Ala Leu Ala Glu 20 25 His Gln Lys Thr His Glu Lys Glu Lys Ser Tyr Ala Leu Gly Gly 35 40 Ala Arg Gly Pro Gln Pro Ser Thr Arg Glu Pro Arg Arg Gly Leu 50 55 Gly Arg Ala Val Pro Gln Arg Ala Trp Arg Ala Arg Leu Pro Pro 65 70 His Pro Gln Arg Arg Gly Glu Pro Leu Cys Cys Pro Val Pro 80 Glu Gly Pro Leu Cys Arg Pro 95

<210> 32 <211> 812 <212> PRT <213> Homo sapiens

<220> <221> misc\_feature <223> Incyte cione 2076520CD1

<400> 32 Met Ile Glu Pro Asp Gln Cys Phe Cys Arg Phe Asp Leu Thr Gly Thr Cys Asn Asp Asp Cys Gln Trp Gln His Ile Gln Asp Tyr Thr Leu Ser Arg Lys Gln Leu Phe Gln Asp Ile Leu Ser Tyr Asn Leu Ser Leu Iie Gly Cys Ala Glu Thr Ser Thr Asn Glu Glu Ile Thr Ala Ser Ala Glu Lys Tyr Val Glu Lys Leu Phe Gly Val Asn Lys Asp Arg Met Ser Met Asp Gln Met Ala Val Leu Leu Val Ser Asn Ile Asn Glu Ser Lys Gly His Thr Pro Pro Phe Thr Thr Tyr Lys Asp Lys Arg Lys Trp Lys Pro Lys Phe Trp Arg Lys Pro Ile Ser Asp Asn Ser Phe Ser Ser Asp Glu Glu Gln Ser Thr Gly Pro Ile Lys Tyr Ala Phe Gln Pro Glu Asn Gln Ile Asn Val Pro Ala Leu Asp Thr Val Val Thr Pro Asp Asp Val Arg Tyr Phe Thr Asn Glu Thr Asp Asp Ile Ala Asn Leu Glu Ala Ser Val Leu Glu Asn Pro Ser His Val Gln Leu Trp Leu Lys Leu Ala Tyr Lys Tyr Leu Asn Gln Asn Glu Gly Glu Cys Ser Glu Ser Leu Asp Ser Ala Leu Asn Val Leu Ala Arg Ala Leu Glu Asn Asn Lys Asp Asn Pro Glu Ile Trp Cys His Tyr Leu Arg Leu Phe Ser Lys Arg Gly Thr Lys Asp Glu Val Gln Glu Met Cys Glu Thr Ala Val Glu Tyr Ala Pro Asp Tyr Gln Ser Phe Trp Thr Phe Leu His Leu Glu Ser Thr Phe Glu Glu Lys Asp Tyr Val Cys Glu Arg Met Leu Glu Phe Leu Met Gly Ala Ala Lys Gln Glu Thr Ser Asn Ile Leu Ser Phe Gln Leu Leu Glu Ala Leu Leu Phe Arg Val Gln Leu His Ile Phe Thr Gly Arg Cys Gln Ser Ala Leu Ala Ile Leu Gln Asn Ala Leu Lys Ser Ala Asn Asp Gly Ile Val Ala Glu Tyr Leu Lys Thr Ser Asp Arg Cys Leu Ala Trp Leu Ala Tyr Ile His Leu Ile Glu Phe Asn Ile Leu Pro Ser Lys Phe Tyr Asp Pro Ser Asn Asp Asn Pro Ser Arg Ile Val Asn Thr Glu Ser Phe Val Met Pro Trp Gln Ala Val Gln Asp Val Lys Thr Asn Pro Asp Met Leu Leu Ala Val Phe Glu Asp Ala Val Lys Ala Cys Thr Asp Glu Ser Leu Ala Val Glu Glu Arg Ile Glu Ala Cys Leu Pro Leu Tyr Thr Asn Met Ile Ala Leu His Gln Leu Leu Glu Arg Tyr Glu Ala Ala Met Glu Leu Cys Lys Ser

```
440
                                                          450
                                     445
Leu Leu Glu Ser Cys Pro Ile Asn Cys Gln Leu Leu Glu Ala Leu
                455
                                     460
Val Ala Leu Tyr Leu Gln Thr Asn Gln His Asp Lys Ala Arg Ala
                470
                                     475
Val Trp Leu Thr Ala Phe Glu Lys Asn Pro Gln Asn Ala Glu Val
                485
                                     490
Phe Tyr His Met Cys Lys Phe Phe Ile Leu Gln Asn Arg Gly Asp
                                     505
                500
                                                          510
Asn Leu Leu Pro Phe Leu Arg Lys Phe Ile Ala Ser Phe Phe Lys
                515
                                     520
                                                          525
Pro Gly Phe Glu Lys Tyr Asn Asn Leu Asp Leu Phe Arg Tyr Leu
                530
                                     535
Leu Asn Ile Pro Gly Pro Ile Asp Ile Pro Ser Arg Leu Cys Lys
                545
                                     550
Gly Asn Phe Asp Asp Met Phe Asn His Gln Val Pro Tyr Leu
                560
                                     565
                                                          570
Trp Leu Ile Tyr Cys Leu Cys His Pro Leu Gln Ser Ser Ile Lys
                575
                                     580
                                                          585
Glu Thr Val Glu Ala Tyr Glu Ala Ala Leu Gly Val Ala Met Arg
                590
                                     595
                                                          600
Cys Asp Ile Val Gln Lys Ile Trp Met Asp Tyr Leu Val Phe Ala
                605
                                     610
                                                          615
Asn Asn Arg Ala Ala Gly Ser Arg Asn Lys Val Gln Glu Phe Arg
                620
                                     625
Phe Phe Thr Asp Leu Val Asn Arg Cys Leu Val Thr Val Pro Ala
                635
                                     640
                                                          645
Arg Tyr Pro Ile Pro Phe Ser Ser Ala Asp Tyr Trp Ser Asn Tyr
                650
                                     655
Glu Fhe His Asn Arg Val Ile Phe Phe Tyr Leu Ser Cys Val Pro
                665
                                     670
                                                          675
Lys Thr Gln His Ser Lys Thr Leu Glu Arg Phe Cys Ser Val Met
                680
                                     685
                                                          690
Pro Ala Asn Ser Gly Leu Ala Leu Arg Leu Leu Gln His Glu Trp
                695
                                     700
                                                          705
Glu Glu Ser Asn Val Gln Ile Leu Lys Leu Gln Ala Lys Met Phe
                710
                                     715
Thr Tyr Asn Ile Pro Thr Cys Leu Ala Thr Trp Lys Ile Ala Ile
                725
                                     730
                                                          735
Ala Ala Glu Ile Val Leu Lys Gly Gln Arg Glu Val His Arg Leu
                740
                                     745
Tyr Gln Arg Ala Leu Gln Lys Leu Pro Leu Cys Ala Ser Leu Trp
                755
                                     760
                                                          765
Lys Asp Gln Leu Leu Phe Glu Ala Ser Glu Gly Gly Lys Thr Asp
                770
                                     775
Asn Leu Arg Lys Leu Val Ser Lys Cys Gln Glu Ile Gly Val Ser
                785
                                     790
                                                          795
Leu Asn Glu Leu Leu Asn Leu Asn Ser Asn Lys Thr Glu Ser Lys
                800
                                     805
Asn His
```

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<210> 33

<211> 392

<212> FRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte clone 2291241CD1

<400> 33
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Met Asp Ala Leu Val Glu Asp Asp Ile Cys Ile Leu Asn His Glu

```
Lys Ala His Lys Arg Asp Thr Val Thr Pro Val Ser Ile Tyr Ser
                                      25
Gly Asp Glu Ser Val Ala Ser His Phe Ala Leu Val Thr Ala Tyr
                                                           45
Glu Asp Ile Lys Lys Arg Leu Lys Asp Ser Glu Lys Glu Asn Ser
                  50
Leu Leu Lys Lys Arg Ile Arg Phe Leu Glu Glu Lys Leu Ile Ala
                  65
                                      70
Arg Phe Glu Glu Glu Thr Ser Ser Val Gly Arg Glu Gln Val Asn
                  80
                                      85
Lys Ala Tyr His Ala Tyr Arg Glu Val Cys Ile Asp Arg Asp Asn
                  95
                                     100
Leu Lys Ser Lys Leu Asp Lys Met Asn Lys Asp Asn Ser Glu Ser
                 110
                                     115
Leu Lys Val Leu Asn Glu Gln Leu Gln Ser Lys Glu Val Glu Leu
                 125
                                     130
Leu Gln Leu Arg Thr Glu Val Glu Thr Gln Gln Val Met Arg Asn
                 140
                                     145
Leu Asn Pro Pro Ser Ser Asn Trp Glu Val Glu Lys Leu Ser Cys
                 155
                                     160
Asp Leu Lys Ile His Gly Leu Glu Gln Glu Leu Glu Leu Met Arg
                 170
                                     175
                                                          180
Lys Glu Cys Ser Asp Leu Lys Ile Glu Leu Gln Lys Ala Lys Gln
                185
                                     190
Thr Asp Pro Tyr Gln Glu Asp Asn Leu Lys Ser Arg Asp Leu Gln
                200
                                     205
Lys Leu Ser Ile Ser Ser Asp Asn Met Gln His Ala Tyr Trp Glu
                215
                                     220
Leu Lys Arg Glu Met Ser Asn Leu His Leu Val Thr Gln Val Gln
                230
                                     235
Ala Glu Leu Leu Arg Lys Leu Lys Thr Ser Thr Ala Ile Lys Lys
                245
                                     250
Ala Cys Ala Pro Val Gly Cys Ser Glu Asp Leu Gly Arg Asp Ser
                260
                                     265
Thr Lys Leu His Leu Met Asn Phe Thr Ala Thr Tyr Thr Arg His
                275
                                     280
Pro Pro Leu Leu Pro Asn Gly Lys Ala Leu Cys His Thr Thr Ser
                290
                                     295
                                                          300
Ser Pro Leu Pro Gly Asp Val Lys Val Leu Ser Glu Lys Ala Ile
                305
                                     310
                                                          315
Leu Gln Ser Trp Thr Asp Asn Glu Arg Ser Ile Pro Asn Asp Gly
                320
                                     325
                                                          330
Thr Cys Phe Gln Glu His Ser Ser Tyr Gly Arg Asn Ser Leu Glu
                335
                                     340
Asp Asn Ser Trp Val Phe Pro Ser Pro Pro Lys Ser Ser Glu Thr
                350
                                     355
Ala Phe Gly Glu Thr Lys Thr Lys Thr Leu Pro Leu Pro Asn Leu
                365
                                     370
                                                         375
Pro Pro Leu His Tyr Leu Asp Gln His Asn Gln Asn Cys Leu Tyr
                380
                                     385
                                                         390
Lys Asn
```

<sup>&</sup>lt;210> 34

<sup>&</sup>lt;211> 60

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

<sup>&</sup>lt;220>

<sup>&</sup>lt;221> misc\_feature

<sup>&</sup>lt;223> Incyte clone 2329692CD1

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<400> 34
Met Ile Tyr Phe Phe Ile Ile Ile Val Glu Tyr Phe Tyr Gly Lys
                                     10
Ile Phe Val Val Leu Ile Ile Pro Ile Lys Ile Met Pro Asn Thr
                                      25
                                                           30
Lys Tyr Glu Phe Tyr Asp Val His Phe Val Leu Gly Ile Lys Arg
                 35
                                      40
Lys Lys His Thr Ser Trp Lys Ser Val Ser Cys Phe Leu Leu
<210> 35
<211> 209
<212> PRT
<213> Homo sapiens
<220>
<221> misc feature
<223> Incyte clone 2474110CD1
<400> 35
Met Asp Pro Ser Asp Ile Tyr Ala Val Ile Gln Ile Pro Gly Ser
 1
                                     1.0
Arg Glu Phe Asp Val Ser Phe Arg Ser Ala Glu Lys Leu Ala Leu
                 20
                                                          30
Phe Leu Arg Val Tyr Glu Glu Lys Arg Glu Gln Glu Asp Cys Trp
                 35
                                      40
Glu Asn Phe Val Val Leu Gly Arg Ser Lys Ser Ser Leu Lys Thr
                 50
                                      55
Leu Phe Ile Leu Phe Arg Asn Glu Thr Val Asp Val Glu Asp Ile
                 65
                                     70
Val Thr Trp Leu Lys Arg His Cys Asp Val Leu Ala Val Pro Val
                 80
                                     85
                                                          90
Lys Val Thr Asp Arg Phe Gly Ile Trp Thr Gly Glu Tyr Lys Cys
                 95
                                    100
                                                         105
Glu Ile Glu Leu Arg Gln Gly Glu Gly Gly Val Arg His Leu Pro
                110
                                    115
Gly Ala Phe Phe Leu Gly Ala Glu Arg Gly Tyr Ser Trp Tyr Lys
                                    130
                                                         135
Gly Gln Pro Lys Thr Cys Phe Lys Cys Gly Ser Arg Thr His Met
                140
                                     145
Ser Gly Ser Cys Thr Gln Asp Arg Cys Phe Arg Cys Arg Glu Glu
                155
                                     160
Gly His Leu Ser Pro Tyr Cys Arg Lys Gly Ile Val Cys Asn Leu
                170
                                    175
Cys Gly Lys Arg Gly His Ala Phe Ala Gln Cys Pro Lys Ala Val
                185
                                    190
His Asn Ser Val Ala Ala Gln Leu Thr Gly Val Ala Gly His
                200
                                    205
<210> 36
<211> 257
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<211> 257
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 2495790CD1
<400> 36
Met Val Gly Ala Gly Ile Ser Thr Pro Ser Gly Ile Pro Asp Phe
```

```
Arg Ser Pro Gly Ser Gly Leu Tyr Ser Asn Leu Gln Gln Tyr Asp
                                      25
Leu Pro Tyr Pro Glu Ala Ile Phe Glu Leu Pro Phe Phe His
                                      40
Asn Pro Lys Pro Phe Phe Thr Leu Ala Lys Glu Leu Tyr Pro Gly
                  50
                                      55
                                                           60
Asn Tyr Lys Pro Asn Val Thr His Tyr Phe Leu Arg Leu Leu His
                  65
                                      70
Asp Lys Gly Leu Leu Leu Arg Leu Tyr Thr Gln Asn Ile Asp Gly
                 80
                                      85
                                                           90
Leu Glu Arg Val Ser Gly Ile Pro Ala Ser Lys Leu Val Glu Ala
                 95
                                     100
His Gly Thr Phe Ala Ser Ala Thr Cys Thr Val Cys Gln Arg Pro
                110
                                     115
Phe Pro Gly Glu Asp Ile Arg Ala Asp Val Met Ala Asp Arg Val
                125
                                     130
                                                         135
Pro Arg Cys Pro Val Cys Thr Gly Val Val Lys Pro Asp Ile Val
                140
                                     145
                                                          150
Phe Phe Gly Glu Pro Leu Pro Gln Arg Phe Leu Leu His Val Val
                155
                                     160
                                                          165
Asp Fhe Pro Met Ala Asp Leu Leu Leu Ile Leu Gly Thr Ser Leu
                                     175
Glu Val Glu Pro Phe Ala Ser Leu Thr Glu Ala Val Arg Ser Ser
                185
                                     190
Val Pro Arg Leu Leu Ile Asn Arg Asp Leu Val Gly Pro Leu Ala
                200
                                     205
                                                         210
Trp His Pro Arg Ser Arg Asp Val Ala Gln Leu Gly Asp Val Val
                215
                                     220
His Gly Val Glu Ser Leu Val Glu Leu Leu Gly Trp Thr Glu Glu
                230
                                     235
Met Arg Asp Leu Val Gln Arg Glu Thr Gly Lys Leu Asp Gly Pro
                245
                                     250
Asp Lvs
```

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<210> 37
<211> 138
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 2661254CD1
<400> 37
Met Ala Thr Lys Arg Leu Phe Gly Ala Thr Arg Thr Trp Ala Gly
Trp Gly Ala Trp Glu Leu Leu Asn Pro Ala Thr Ser Gly Arg Leu
                 20
Leu Ala Arg Asp Tyr Ala Lys Lys Pro Val Met Lys Gly Ala Lys
                 35
```

50

80

95

110

10

25

40

55

70

85

100

115

130

60

105

120

135

Ser Gly Lys Gly Ala Val Thr Ser Glu Ala Leu Lys Asp Pro Asp

Val Cys Thr Asp Pro Val Gln Leu Thr Thr Tyr Ala Met Gly Val

Asn Ile Tyr Lys Glu Gly Gln Asp Val Pro Leu Lys Pro Asp Ala

Glu Tyr Pro Glu Trp Leu Phe Glu Met Asn Leu Gly Pro Pro Lys

Thr Leu Glu Glu Leu Asp Pro Glu Ser Arg Glu Tyr Trp Arg Arg

Leu Arg Lys Gln Asn Ile Trp Arg His Asn Arg Leu Ser Lys Asn

<210> 38

Lys Arg Leu

```
<211> 999
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 2674047CD1
<400> 38
Met Gly Pro Ser Arg Leu Arg Leu Gly Phe Phe Xaa Lys Arg Gly
                                      10
                                                           15
Cys Ser Arg Ala Met Val Glu Ile Glu Leu Phe Arg Ala Ser Gly
                 20
                                      25
                                                           30
Asn Leu Val Ile Thr Arg Glu Ile Asp Val Ala Lys Asn Gln Ser
                 35
                                      40
                                                           45
Phe Trp Phe Ile Asn Lys Lys Ser Thr Thr Gln Xaa Ile Val Glu
                 50
                                      55
Glu Lys Val Ala Ala Leu Asn Ile Gln Val Gly Asn Leu Cys Gln
                                                           7.5
Phe Leu Pro Glm Asp Lys Val Gly Glu Phe Ala Lys Leu Ser Lys
                 80
Ile Glu Leu Leu Glu Ala Thr Glu Lys Ser Ile Gly Pro Pro Glu
                 95
                                     100
Met His Lys Tyr His Cys Glu Leu Lys Asn Leu Arg Glu Lys Glu
                110
                                     115
Lys Gln Leu Glu Thr Ser Cys Lys Glu Lys Thr Glu Tyr Leu Gln
                125
                                     130
                                                          135
Lys Met Val Gln Arg Asn Glu Arg Tyr Lys Gln Asp Val Glu Arg
                140
                                     145
Phe Tyr Glu Arg Lys Arg His Leu Asp Leu Ile Glu Met Leu Glu
                155
                                     160
                                                          165
Ala Lys Arg Pro Trp Val Glu Tyr Glu Asn Val Arg Gln Glu Tyr
                170
                                     175
                                                          180
Glu Glu Val Lys Leu Val Arg Asp Arg Val Lys Glu Glu Val Arg
                185
                                     190
                                                          195
Lys Leu Lys Glu Gly Gln Ile Pro Ile Thr Cys Arg Ile Glu Glu
                200
                                     205
                                                          210
Met Glu Asn Glu Arg His Asn Leu Glu Ala Arg Ile Lys Glu Lys
                215
                                     220
Ala Thr Asp Ile Lys Glu Ala Ser Gln Lys Cys Lys Gln Lys Gln
                230
                                     235
Asp Val Ile Glu Arg Lys Asp Lys His Ile Glu Glu Leu Gln Gln
                245
                                     250
                                                          255
Ala Leu Ile Val Lys Gln Asn Glu Glu Leu Asp Arg Gln Arg Arg
                260
                                     265
Ile Gly Asn Thr Arg Lys Met Ile Glu Asp Leu Gln Asn Glu Leu
                275
                                     280
                                                          285
Lys Thr Thr Glu Asn Cys Glu Asn Leu Gln Pro Gln Ile Asp Ala
                290
                                     295
                                                          300
Ile Thr Asn Asp Leu Arg Arg Ile Gln Asp Glu Lys Ala Leu Cys
                305
                                     310
                                                          315
Glu Gly Glu Ile Ile Asp Lys Arg Arg Glu Arg Glu Thr Leu Glu
                320
                                     325
                                                          330
Lys Glu Lys Lys Ser Val Asp Asp His Ile Val Arg Phe Asp Asn
                335
                                     340
                                                          345
Leu Met Asn Gln Lys Glu Asp Lys Leu Arg Gln Arg Phe Arg Asp
                350
                                     355
                                                          360
Thr Tyr Asp Ala Val Leu Trp Leu Arg Asn Asn Arg Asp Lys Phe
                365
                                     370
```

Lys	s Gli	n Arc	y Val	Cys 380	s Glu	Pro	o Ile	e Mei	t Let 385	Thr	: Ile	Asn	Met	Lys
Asp	) Ası	ı Lys	. Asr	n Ala 395	Lys	туг	: Ile	e Glı	Asr 400	His	Ile	Pro	Sei	390 Asn
Asp	) Let	ı Arç	, Ala	Phe 410	val	Phe	Glu	ı Sei	Gln 415	Glu	Asp	Met	Glu	405 Val 420
Phe	e Leu	ı Lys	Glu	Val 425	Arg	Asp	Asr	Lys	Lys 430	Leu	Arg	Val	Asr	420 Ala 435
Val	. Ile	e Ala	Pro	Lys 440	Ser	Ser	Tyr	Ala	445 445	Lys	Ala	Pro	Ser	Arg
Ser	Leu	Asn	Glu	Leu 455	Lys	Gln	Tyr	: Gl	Phe 460	Phe	Ser	Tyr	Leu	450 Arg 465
Glu	Leu	Phe	Asp	Ala 470	Pro	Asp	Pro	Val	. Met 475	Ser	Tyr	Leu	Cys	Cys 480
				485					. Gly 490	Thr				Arg
				500					Thr 505					Ile 510
				212					Lys 520					Ser
				530	Ser				Leu 535					Phe
				245					Arg 550					Glu 555
				560	His				Gln 565					Gly 570
				2/5	Glu				580					Asp
				590	Lys				595					Thr
				605	Glu				610					Ser
				620	Gln				625					630
				635	Lys				640					615
				650	Leu				655					660
				665	Val				670					675
				680	Lys				685					690
		Leu		695					Phe 700					705
				/10	Leu				715					720
				125	Asn				730					725
				740	Val				745					750
				/55	Phe				760					765
				//0	Thr				775					780
				785	Thr				790					705
				800	Leu				805					810
				812	Arg				ጸጋበ					025
				830	Lys				835					240
			- 116	845	Ser	261	ne t	GTU	850	ATA	GIĀ	GLu	Val	Asp 855

```
Leu His Thr Glu Asn Glu Glu Asp Tyr Asp Lys Tyr Gly Ile Arg
                                     865
Ile Arg Val Lys Phe Arg Ser Ser Thr Gln Leu His Glu Leu Thr
                875
                                     880
Pro His His Gln Ser Gly Gly Glu Arg Ser Val Ser Thr Met Leu
                890
                                     895
Tyr Leu Met Ala Leu Gln Glu Leu Asn Arg Cys Pro Phe Arg Val
                905
                                     910
                                                         915
Val Asp Glu Ile Asn Gln Gly Met Asp Pro Ile Asn Glu Arg Arg
                920
                                     925
                                                         930
Val Phe Glu Met Val Val Asn Thr Ala Cys Lys Glu Asn Thr Ser
                935
                                     940
                                                         945
Gln Tyr Phe Phe Ile Thr Pro Lys Leu Leu Gln Asn Leu Pro Tyr
                950
                                     955
                                                         960
Ser Glu Lys Met Thr Val Leu Phe Val Tyr Asn Gly Pro His Met
                965
                                    970
Leu Glu Pro Asn Thr Trp Asn Leu Lys Ala Phe Gln Arg Arg
                980
                                    985
                                                         990
Arg Arg Ile Thr Phe Thr Gln Pro Ser
                995
```

<210> 39 <211> 377 <212> PRT <213> Homo sapiens

<221> misc feature <223> Incyte clone 2762174CD1

<400> 39 Met Ala Glu Leu Glu Ser His Pro Cys Asp Ile Cys Gly Pro Ile Leu Lys Asp Thr Leu His Leu Ala Lys Tyr His Gly Gly Lys Ala Arg Gln Lys Pro Tyr Leu Cys Gly Ala Cys Gly Lys Gln Phe Trp Phe Ser Thr Asp Phe Asp Gln His Gln Asn Gln Pro Asn Gly Gly Lys Leu Phe Pro Arg Lys Glu Gly Arg Asp Ser Val Lys Ser Cys Arg Val His Val Pro Glu Lys Thr Leu Thr Cys Gly Lys Gly Arg Arg Asp Phe Ser Ala Thr Ser Gly Leu Leu Gln His Gln Ala Ser Leu Ser Ser Met Lys Pro His Lys Ser Thr Lys Leu Val Ser Gly Phe Leu Met Gly Gln Arg Tyr His Arg Cys Gly Glu Cys Gly Lys Ala Phe Thr Arg Lys Asp Thr Leu Ala Arg His Gln Arg Ile His Thr Gly Glu Arg Pro Tyr Glu Cys Asn Glu Cys Gly Lys Phe Phe Ser Gln Ser Tyr Asp Leu Phe Lys His Gln Thr Val His Thr Gly Glu Arg Pro Tyr Glu Cys Ser Glu Cys Gly Lys Phe Phe Arg Gln Ile Ser Gly Leu Ile Glu His Arg Arg Val His Thr Gly Glu Arg Leu Tyr Gln Cys Gly Lys Cys Gly Lys Phe Phe Ser Ser Lys Ser Asn Leu Ile Arg His Gln Glu Val His Thr Gly Ala Arg Pro Tyr

```
Val Cys Ser Glu Cys Gly Lys Glu Phe Ser Arg Lys His Thr Leu
                                     250
Val Leu His Gln Arg Thr His Thr Gly Glu Arg Pro Tyr Glu Cys
                260
                                     265
                                                          270
Ser Glu Cys Gly Lys Ala Phe Ser Gln Ser Ser His Leu Asn Val
                275
                                     280
                                                          285
His Trp Arg Ile His Ser Ser Asp Tyr Glu Cys Ser Arg Cys Gly
                290
                                     295
                                                          300
Lys Ala Phe Ser Cys Ile Ser Lys Leu Ile Gln His Gln Lys Val
                305
                                     310
His Ser Gly Glu Lys Pro Tyr Glu Cys Ser Lys Cys Gly Lys Ala
                320
                                     325
Phe Thr Gln Arg Pro Asn Leu Ile Arg His Trp Lys Val His Thr
                335
                                     340
Gly Glu Arg Pro Tyr Val Cys Ser Glu Cys Gly Arg Glu Phe Ile
                350
                                     355
Arg Lys Gln Thr Leu Val Leu His Gln Arg Val His Ala Gly Glu
                365
                                     370
                                                          375
Lys Leu
```

<210> 40 <211> 324 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte clone 2765991CD1 <400> 40

Met Asp Phe Pro Lys His Asn Gln Ile Ile Thr Glu Glu Thr Gly Ser Ala Val Glu Pro Ser Asp Glu Ile Lys Arg Ala Ser Gly Asp Val Gln Thr Met Lys Ile Ser Ser Val Pro Asn Ser Leu Ser Lys Arg Asn Val Ser Leu Thr Arg Ser His Ser Val Gly Gly Pro Leu Gln Asn Ile Asp Phe Thr Gln Arg Pro Phe His Gly Ile Ser Thr Val Ser Leu Pro Gly Ser Leu Gln Glu Val Val Asp Pro Leu Gly Lys Arg Pro Asn Pro Pro Pro Val Ser Val Pro Tyr Leu Ser Pro Leu Val Leu Arg Lys Glu Leu Glu Ser Leu Leu Glu Asn Glu Gly Asp Gln Val Ile His Thr Ser Ser Phe Ile Asn Gln His Pro Ile Ile Phe Trp Asn Leu Val Trp Tyr Phe Arg Arg Leu Asp Leu Pro Ser Asn Leu Pro Gly Leu Ile Leu Thr Ser Glu His Cys Asn Glu Gly Val Gln Leu Pro Leu Ser Ser Leu Ser Gln Asp Ser Lys Leu Val Tyr Ile Arg Leu Leu Trp Asp Asn Ile Asn Leu His Gln Glu Pro Arg Glu Pro Leu Tyr Val Ser Trp Arg Asn Phe Asn Ser Glu Lys Lys Ser Ser Leu Leu Ser Glu Glu Gln Gln Glu Thr Ser Thr Leu Val Glu Thr Ile Arg Gln Ser Ile Gln His Asn Asn Val Leu 

```
Lys Pro Ile Asn Leu Leu Ser Gln Gln Met Lys Pro Gly Met Lys
                245
                                     250
                                                          255
Arg Gln Arg Ser Leu Tyr Arg Glu Ile Leu Phe Leu Ser Leu Val
                                     265
                                                          270
Ser Leu Gly Arg Glu Asn Ile Asp Ile Glu Ala Phe Asp Asn Glu
                275
                                     280
                                                          285
Tyr Gly Ile Ala Tyr Asn Ser Leu Ser Ser Glu Ile Leu Glu Arg
                290
                                     295
                                                          300
Leu Gln Lys Ile Asp Ala Pro Pro Ser Ala Ser Val Glu Trp Cys
                305
                                     310
                                                          315
Arg Lys Cys Phe Gly Ala Pro Leu Ile
                320
```

Ser Thr Leu Glu Lys Cys Asn Arg Ser Glu Ala Thr Trp Gly Arg Val Pro Ser Asp Pro Phe Thr Gly Val Ala Phe Thr Pro His Ser Gln Pro Leu Pro His Pro Ser Leu Lys Ala Arg Ile Asp His Phe Leu Leu Gln His Ser Ile Pro Gly Cys His Leu Leu Gly Arg Ala Gln Thr Ala Leu Ala Val Ile Pro Ser Ser Ile Val Leu Pro Ser Gln Lys Arg Lys Ile Glu Gln Ala Glu His Val Pro Asp Ser Asn Phe Gly Val Asn Ala Ser Cys Phe Ser Ala Thr Ser Pro Leu Val Leu Pro Thr Thr Ser Glu His Thr Ala Lys Lys Met Lys Ala Thr Asn Glu Pro Ser Leu Thr His Met Asp Cys Ser Thr Gly Pro Leu Ser His Glu Gln Lys Leu Ser Gln Ser Leu Glu Ile Ala Leu Ala Ser Thr Leu Gly Ser Met Pro Ser Phe Thr Ala Arg Leu Thr Arg Gly Gln Leu Gln His Leu Gly Thr Arg Gly Ser Asn Thr Ser Trp Arg Pro Gly Thr Gly Ser Glu Gln Pro Gly Ser Ile Leu Gly Pro Glu Cys Ala Ser Cys Lys Arg Val Phe Ser Pro Tyr Phe Lys Lys Glu Pro Val Tyr Gln Leu Pro Cys Gly His Leu Leu Cys Arg Pro Cys Leu Gly Glu Lys Gln Arg Ser Leu Pro Met Thr Cys Thr Ala Cys Gln Arg Pro Val Ala Ser Gln Asp Val Leu Arg Val His Phe

<210> 42 <211> 252 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte clone 2918375CD1 <400> 42 Met Leu Arg Lys Gly Ile Cys Glu Tyr His Glu Lys Asn Tyr Ala 10 Ala Ala Leu Glu Thr Phe Thr Glu Gly Gln Lys Leu Asp Ser Ala 20 25 30 Asp Ala Asn Phe Ser Val Trp Ile Lys Arg Cys Gln Glu Ala Gln 35 40 45 Asn Gly Ser Glu Ser Glu Val Trp Thr His Gln Ser Lys Ile Lys 50 55 60 Tyr Asp Trp Tyr Gln Thr Glu Ser Gln Val Val Ile Thr Leu Met 65 70 75 Ile Lys Asn Val Gln Lys Asn Asp Val Asn Val Glu Phe Ser Glu 80 85 90 Lys Glu Leu Ser Ala Leu Val Lys Leu Pro Ser Gly Glu Asp Tyr 95 100 105 Asn Leu Lys Leu Glu Leu Leu His Pro Ile Ile Pro Glu Gln Ser 110 115 120 Thr Phe Lys Val Leu Ser Thr Lys Ile Glu Ile Lys Leu Lys Lys 125 130 135 Pro Glu Ala Val Arg Trp Glu Lys Leu Glu Gly Gln Gly Asp Val 140 145 150 Pro Thr Pro Lys Gln Phe Val Ala Asp Val Lys Asn Leu Tyr Pro 155 160 165 Ser Ser Ser Pro Tyr Thr Arg Asn Trp Asp Lys Leu Val Gly Glu 170 175 Ile Lys Glu Glu Glu Lys Asn Glu Lys Leu Glu Gly Asp Ala Ala 185 190 195 Leu Asn Arg Leu Phe Gln Gln Ile Tyr Ser Asp Gly Ser Asp Glu 200 205 210 Val Lys Arg Ala Met Asn Lys Ser Phe Met Glu Ser Gly Gly Thr 215 220 225 Val Leu Ser Thr Asn Trp Ser Asp Val Gly Lys Arg Lys Val Glu 230 235 Ile Asn Pro Pro Asp Asp Met Glu Trp Lys Lys Tyr 245 <210> 43 <211> 228 <212> PRT

```
<213> Homo sapiens
<220>
<221> misc feature
<223> Incyte clone 3149729CD1
<400> 43
Met Thr Met Gly Asp Lys Lys Ser Pro Thr Arg Pro Lys Arg Gln
 1
                                      10
Ala Lys Pro Ala Ala Asp Glu Gly Phe Trp Asp Cys Ser Val Cys
                 20
                                      25
                                                           30
Thr Phe Arg Asn Ser Ala Glu Ala Phe Lys Cys Ser Ile Cys Asp
                 35
                                      40
Val Arg Lys Gly Thr Ser Thr Arg Lys Pro Arg Ile Asn Ser Gln
                 50
```

```
Leu Val Ala Gln Gln Val Ala Gln Gln Tyr Ala Thr Pro Pro
                 65
Pro Lys Lys Glu Lys Glu Lys Val Glu Lys Gln Asp Lys Glu
                                                          90
                                     85
Lys Pro Glu Lys Asp Lys Glu Ile Ser Pro Ser Val Thr Lys Lys
                 95
                                    100
                                                         105
Asn Thr Asn Lys Lys Thr Lys Pro Lys Ser Asp Ile Leu Lys Asp
                110
                                    115
                                                         120
Pro Pro Ser Glu Ala Asn Ser Ile Gln Ser Ala Asn Ala Thr Thr
                125
                                    130
Lys Thr Ser Glu Thr Asn His Thr Ser Arg Pro Arg Leu Lys Asn
                140
                                    145
Val Asp Arg Ser Thr Ala Gln Gln Leu Ala Val Thr Val Gly Asn
                155
                                    160
Val Thr Val Ile Ile Thr Asp Phe Lys Glu Lys Thr Arg Ser Ser
                170
                                    175
Ser Thr Ser Ser Ser Thr Val Thr Ser Ser Ala Gly Ser Glu Gln
                185
                                    190
                                                         195
Gln Asn Gln Ser Ser Ser Gly Ser Glu Ser Thr Asp Lys Gly Ser
                200
                                    205
Ser Arg Ser Ser Thr Pro Lys Gly Asp Met Ser Ala Val Asn Asp
                                    220
Glu Ser Phe
```

```
<210> 44
<211> 117
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 3705895CD1
<400> 44
Met Ala Ala Ala Ala Ala Gly Ser Gly Thr Pro Arg Glu Glu
 1
                                     10
Glu Gly Pro Ala Gly Glu Ala Ala Ser Gln Pro Gln Ala Pro
Thr Ser Val Pro Gly Ala Arg Leu Ser Arg Leu Pro Leu Ala Arg
                 35
                                     40
                                                          4.5
Val Lys Ala Leu Val Lys Ala Asp Pro Asp Val Thr Leu Ala Gly
                 50
                                     55
Gln Glu Ala Ile Phe Ile Leu Ala Arg Ala Ala Glu Leu Phe Val
                 65
                                     70
                                                          75
Glu Thr Ile Ala Lys Asp Ala Tyr Cys Cys Ala Gln Gln Gly Lys
                 80
                                     85
Arg Lys Thr Leu Gln Arg Arg Asp Leu Asp Asn Ala Ile Glu Ala
                 95.
                                    100
                                                         105
Val Asp Glu Phe Ala Phe Leu Glu Gly Thr Leu Asp
                110
                                    115
```

```
<210> 45
<211> 252
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 003256CD1
```

```
<400> 45
Met Thr Pro Lys Leu Gly Arg Gly Val Leu Glu Gly Asp Asp Val
                                      10
Leu Phe Tyr Asp Glu Ser Pro Pro Pro Arg Pro Lys Leu Ser Ala
                 20
                                      25
                                                           30
Leu Ala Glu Ala Lys Lys Leu Ala Ala Ile Thr Lys Leu Arg Ala
                                      40
Lys Gly Gln Val Leu Thr Lys Thr Asn Pro Asn Ser Ile Lys Lys
                 50
                                      55
                                                           60
Lys Gln Lys Asp Pro Gln Asp Ile Leu Glu Val Lys Glu Arg Val
                 65
                                      70
Glu Lys Asn Thr Met Phe Ser Ser Gln Ala Glu Asp Glu Leu Glu
                 80
                                      85
Pro Ala Arg Lys Lys Arg Arg Glu Gln Leu Ala Tyr Leu Glu Ser
                 95
                                     100
                                                          105
Glu Glu Phe Gln Lys Ile Leu Lys Ala Lys Ser Lys His Thr Gly
                110
                                     115
                                                          120
Ile Leu Lys Glu Ala Glu Ala Glu Met Gln Glu Arg Tyr Phe Glu
                125
                                     130
                                                          135
Pro Leu Val Lys Lys Glu Gln Met Glu Glu Lys Met Arg Asn Ile
                140
                                     145
Arg Glu Val Lys Cys Arg Val Val Thr Cys Lys Thr Cys Ala Tyr
                155
                                     160
                                                          165
Thr His Phe Lys Leu Leu Glu Thr Cys Val Ser Glu Gln His Glu
                170
                                     175
                                                          180
Tyr His Trp His Asp Gly Val Lys Arg Phe Phe Lys Cys Pro Cys
                185
                                     190
                                                         195
Gly Asn Arg Ser Ile Ser Leu Asp Arg Leu Pro Asn Lys His Cys
                200
                                     205
Ser Asn Cys Gly Leu Tyr Lys Trp Glu Arg Asp Gly Met Leu Lys
                215
                                     220
                                                         225
Glu Lys Thr Gly Pro Lys Ile Gly Gly Glu Thr Leu Leu Pro Arg
                230
                                     235
                                                         240
Gly Glu Glu His Ala Lys Phe Leu Asn Ser Leu Lys
                245
```

```
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 156986CD1
<400> 46
Met Ala Lys Gly Glu Gly Ala Glu Ser Gly Ser Ala Ala Gly Leu
                                      10
Leu Pro Thr Ser Ile Leu Gln Ser Thr Glu Arg Pro Ala Gln Val
                 20
                                      25
                                                          30
Lys Lys Glu Pro Lys Lys Lys Gln Gln Leu Ser Val Cys Asn
                 35
                                      40
Lys Leu Cys Tyr Ala Leu Gly Gly Ala Pro Tyr Gln Val Thr Gly
                 50
                                      55
                                                          60
Cys Ala Leu Gly Phe Phe Leu Gln Ile Tyr Leu Leu Asp Val Ala
                 65
                                      70
                                                          75
Gln Val Gly Pro Phe Ser Ala Ser Ile Ile Leu Phe Val Gly Arg
                 80
                                      85
                                                          90
Ala Trp Asp Ala Ile Thr Asp Pro Leu Val Gly Leu Cys Ile Ser
                 95
                                     100
Lys Ser Pro Trp Thr Cys Leu Gly Arg Leu Met Pro Trp Ile Ile
                110
                                     115
```

<210> 46 <211> 530

Phe	Ser	Thr	Pro	Leu 125	Ala	Val	Ile	Ala	Tyr 130	Phe	Leu	Ile	Trp	Phe 135
Val	Pro	Asp	Phe		His	Gly	Gln	Thr		Trp	Tyr	Leu	Leu	Phe 150
Tyr	Cys	Leu	Phe		Thr	Met	Val	Thr		Phe	His	Val	Pro	Tyr 165
Ser	Ala	Leu	Thr		Phe	Ile	Ser	Thr		Gln	Thr	Glu	Arg	Asp 180
Ser	Ala	Thr	Ala	Tyr 185	Arg	Met	Thr	Val		Val	Leu	Gly	Thr	
Leu	Gly	Thr	Ala	Ile 200	Gln	Gly	Gln	Ile		Gly	Gln	Ala	Asp	Thr 210
Pro	Cys	Phe	Gln		Leu	Asn	Ser	Ser		Val	Ala	Ser	Gln	Ser 225
Ala	Asn	His	Thr	His 230	Gly	Thr	Thr	Ser	His 235	Arg	Glu	Thr	Gln	Lys 240
Ala	Tyr	Leu	Leu	Ala 245	Ala	Gly	Val	Ile		Суѕ	Ile	Tyr	Ile	Ile 255
Cys	Ala	Val	Ile		Ile	Leu	Gly	Val	Arg 265	Glu	Gln	Arg	Glu	Pro 270
Tyr	Glu	Ala	Gln	Gln 275	Ser	Glu	Pro	Ile		Tyr	Phe	Arg	Gly	Leu 285
Arg	Leu	Val	Met	Ser 290	His	Gly	Pro	Tyr		Lys	Leu	Ile	Thr	Gly 300
Phe	Leu	Phe	Thr	Ser 305	Leu	Ala	Phe	Met		Val	Glu	Gly	Asn	Phe 315
Val	Leu	Phe	Cys	Thr 320	Tyr	Thr	Leu	Gly		Arg	Asn	Glu	Phe	Gln 330
Asn	Leu	Leu	Leu	Ala 335	Ile	Met	Leu	Ser		Thr	Leu	Thr	Ile	Pro
Ile	Trp	Gln	Trp	Phe 350	Leu	Thr	Arg	Phe	Gly 355	Lys	Lys	Thr	Ala	Val 360
Tyr	Val	Gly	Ile	Ser 365	Ser	Ala	Val	Pro	Phe 370	Leu	Ile	Leu	Val	Ala 375
				380				Thr	385					390
				395				Phe	400			_		405
				410				His	415					420
				425				Ser	430					435
				440				Gly	445					450
				455				Gly	460					465
				470				Val	475					480
				485				Phe	490					495
Glu	Glu	Arg	Arg	Arg 500	Gln	Asn	Lys	Lys	Ala 505	Leu	Gln	Ala	Leu	Arg 510
				515	Ser	Gly	Cys	Ser	Glu 520	Thr	Asp	Ser	Thr	Glu 525
Leu	Ala	Ser	Ile	Leu 530										

530

<210> 47 <211> 355 <212> PRT <213> Homo sapiens

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<221> misc_feature
<223> Incyte clone 319415CD1
<400> 47
Met Gly Cys Val Phe Gln Ser Thr Glu Asp Lys Cys Ile Phe Lys
                                      10
Ile Asp Trp Thr Leu Ser Pro Gly Glu His Ala Lys Asp Glu Tyr
                 20
                                      25
                                                          30
Val Leu Tyr Tyr Ser Asn Leu Ser Val Pro Ile Gly Arg Phe
                 35
                                      40
Gln Asn Arg Val His Leu Met Gly Asp Ile Leu Cys Asn Asp Gly
                 50
                                      55
Ser Leu Leu Gln Asp Val Gln Glu Ala Asp Gln Gly Thr Tyr
                 65
                                      70
Ile Cys Glu Ile Arg Leu Lys Gly Glu Ser Gln Val Phe Lys Lys
                 80
                                      85
Ala Val Val Leu His Val Leu Pro Glu Glu Pro Lys Glu Leu Met
                 95
                                     100
Val His Val Gly Gly Leu Ile Gln Met Gly Cys Val Phe Gln Ser
                110
                                     115
                                                         120
Thr Glu Val Lys His Val Thr Lys Val Glu Trp Ile Phe Ser Gly
                                     130
                                                         135
Arg Arg Ala Lys Glu Glu Ile Val Phe Arg Tyr Tyr His Lys Leu
                140
                                     145
Arg Met Ser Val Glu Tyr Ser Gln Ser Trp Gly His Phe Gln Asn
                155
                                     160
Arg Val Asn Leu Val Gly Asp Ile Phe Arg Asn Asp Gly Ser Ile
                170
                                     175
Met Leu Gln Gly Val Arg Glu Ser Asp Gly Gly Asn Tyr Thr Cys
                185
                                     190
                                                         195
Ser Ile His Leu Gly Asn Leu Val Phe Lys Lys Thr Ile Val Leu
                200
                                    205
                                                         210
His Val Ser Pro Glu Glu Pro Arg Thr Leu Val Thr Pro Ala Ala
                215
                                    220
                                                         225
Leu Arg Pro Leu Val Leu Gly Gly Asn Gln Leu Val Ile Ile Val
                230
                                    235
Gly Ile Val Cys Ala Thr Ile Leu Leu Leu Pro Val Leu Ile Leu
                                     250
                                                         255
Ile Val Lys Lys Thr Cys Gly Asn Lys Ser Ser Val Asn Ser Thr
                260
                                     265
Val Leu Val Lys Asn Thr Lys Lys Thr Asn Pro Glu Ile Lys Glu
                275
                                    280
Lys Pro Cys His Phe Glu Arg Cys Glu Gly Glu Lys His Ile Tyr
                290
                                    295
Ser Pro Ile Ile Val Arg Glu Val Ile Glu Glu Glu Pro Ser
                305
                                    310
Glu Lys Ser Glu Ala Thr Tyr Met Thr Met His Pro Val Trp Pro
                320
                                    325
Ser Leu Arg Ser Asp Arg Asn Asn Ser Leu Glu Lys Lys Ser Gly
                335
                                    340
                                                         345
Gly Gly Met Pro Lys Thr Gln Gln Ala Phe
                350
```

```
<210> 48
<211> 136
<212> PRT
<213> Homo sapiens
<220>
<221> misc feature
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<220>

<223> Incyte clone 635581CD1

<210> 49

```
<400> 48
Met Val Gly Gln Thr Glu Asp Asp Thr Ala Gln Gln Leu Val Pro
                                                          15
Thr Cys Gly Met Lys Gly Val Gly Glu Arg Ile Val Glu Tyr Val
Ser Asn Ile Pro Ala Leu Gln Arg Ala Thr Pro Lys Gly Leu Ala
                 35
                                     40
Ser Val Ser Pro Asp Leu Glu His Arg Gln Glu Trp Thr Tyr Ser
                 50
                                     55
Lys Ser Pro Leu Met Gly Lys Gly Thr Arg Leu Glu Ala Ser Glu
                 65
                                     70
Asn Lys Arg Ala Gly Trp Leu Ala Ala Ala Pro Glu Asn Leu Lys
                 80
                                     85
Tyr His Arg Gln Ile Ala Gln Gly Ala Lys Asp Tyr Glu Ile Leu
                 95
                                    100
Lys Lys Glu Thr Asn Lys Phe Ile Leu Arg Ile Tyr Thr His Trp
                110
                                    115
                                                         120
Ser Arg Arg Ser Ile Leu Arg Lys Gly Ser Lys Gly Met Gln Asn
                125
                                    130
Leu
```

<211> 230 <212> PRT <213> Homo sapiens <220> <221> misc feature <223> Incyte clone 921803CD1 <400> 49 Met Lys Leu Ile Val Gly Ile Gly Gly Met Thr Asn Gly Gly Lys 10 Thr Thr Leu Thr Asn Ser Leu Leu Arg Ala Leu Pro Asn Cys Cys 20 30 Val Ile His Gln Asp Asp Phe Phe Lys Pro Gln Asp Gln Ile Ala 35 Val Gly Glu Asp Gly Phe Lys Gln Trp Asp Val Leu Glu Ser Leu 50 55 Asp Met Glu Ala Met Leu Asp Thr Val Gln Ala Trp Leu Ser Ser 65 70 Pro Gln Lys Phe Ala Arg Ala His Gly Val Ser Val Gln Pro Glu 80 85 Ala Ser Asp Thr His Ile Leu Leu Leu Glu Gly Phe Leu Leu Tyr 95 100 Ser Tyr Lys Pro Leu Val Asp Leu Tyr Ser Arg Arg Tyr Phe Leu 110 115 120 Thr Val Pro Tyr Glu Glu Cys Lys Trp Arg Arg Ser Thr Arg Asn 125 130 135 Tyr Thr Val Pro Asp Pro Pro Gly Leu Phe Asp Gly His Val Trp 140 145 Pro Met Tyr Gln Lys Tyr Arg Gln Glu Met Glu Ala Asn Gly Val 155 160 165 Glu Val Val Tyr Leu Asp Gly Met Lys Ser Arg Glu Glu Leu Phe 170 175 Arg Glu Val Leu Glu Asp Ile Gln Asn Ser Leu Leu Asn Arg Ser 185 190 195 Gln Glu Ser Ala Pro Ser Pro Ala Arg Pro Ala Arg Thr Gln Gly 200 205

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Pro Gly Arg Gly Cys Gly His Arg Thr Ala Arg Pro Ala Ala Ser
                215
                                     220
Gln Gln Asp Ser Met
<210> 50
<211> 70
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 1250492CD1
<400> 50
Met Thr Ile Lys Leu Arg Pro Leu Pro Phe Phe Lys Pro Lys Ser
                                      10
Gly Asn Gln Glu Gln Leu His Gly Leu Leu Ala Pro Asp Gln
                 20
Pro Gly Ser Gly Asp Ile Val Ser Leu Phe Gly Asn Cys Arg Pro
                 35
                                      40
Gln Gly Val Gly Leu Ser His Phe Leu Val Leu Pro Thr Phe Pro
                 50
                                      55
Ile Arg Ala Ser Ser Arg Gly Gln Val Cys
<210> 51
<211> 169
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 1427838CD1
<400> 51
Met Leu Ala Phe Ser Glu Met Pro Lys Pro Pro Asp Tyr Ser Glu
 1
                                      10
Leu Ser Asp Ser Leu Thr Leu Ala Val Gly Thr Gly Arg Phe Ser
                 20
                                                          30
Gly Pro Leu His Arg Ala Trp Arg Met Met Asn Phe Arg Gln Arg
                 35
                                      40
Met Gly Trp Ile Gly Val Gly Leu Tyr Leu Leu Ala Ser Ala Ala
                 50
                                      55
Ala Phe Tyr Tyr Val Phe Glu Ile Ser Glu Thr Tyr Asn Arg Leu
                 65
                                      70
                                                          75
Ala Leu Glu His Ile Gln Gln His Pro Glu Glu Pro Leu Glu Gly
                 80
                                      85
Thr Thr Trp Thr His Ser Leu Lys Ala Gln Leu Leu Ser Leu Pro
                 95
                                     100
                                                         105
Phe Trp Val Trp Thr Val Ile Phe Leu Val Pro Tyr Leu Gln Met
                110
                                    115
Phe Leu Phe Leu Tyr Ser Cys Thr Arg Ala Asp Pro Lys Thr Val
                125
                                    130
                                                         135
Gly Tyr Cys Ile Ile Pro Ile Cys Leu Ala Val Ile Cys Asn Arg
                140
                                     145
                                                         150
His Gln Ala Phe Val Lys Ala Ser Asn Gln Ile Ser Arg Leu Gln
                155
                                    160
                                                         165
Leu Ile Asp Thr
```

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<211> 359
<212> PRT
<213> Homo sapiens
<220>
<221> misc feature
<223> Incyte clone 1448258CD1
<400> 52
Met Gly Pro Thr Lys Phe Thr Gln Thr Asn Ile Gly Ile Ile Glu
                                      10
Asn Lys Leu Leu Glu Ala Pro Asp Val Leu Cys Leu Arg Leu Ser
                  20
                                      25
Thr Glu Gln Cys Gln Ala His Glu Glu Lys Gly Ile Glu Glu Leu
                  35
                                      40
Ser Asp Pro Ser Gly Pro Lys Ser Tyr Ser Ile Thr Glu Lys His
                 50
                                      55
Tyr Ala Gln Glu Asp Pro Arg Met Leu Phe Val Ala Ala Val Asp
                 65
                                      70
His Ser Ser Ser Gly Asp Met Ser Leu Leu Pro Ser Ser Asp Pro
                 80
                                      85
Lys Phe Gln Gly Leu Gly Val Val Glu Ser Ala Val Thr Ala Asn
                 95
                                     100
Asn Thr Glu Glu Ser Leu Phe Arg Ile Cys Ser Pro Leu Ser Gly
                110
                                     115
                                                          120
Ala Asn Glu Tyr Ile Ala Ser Thr Asp Thr Leu Lys Thr Glu Glu
                125
                                     130
Val Leu Leu Phe Thr Asp Gln Thr Asp Asp Leu Ala Lys Glu Glu
                140
                                     145
Pro Thr Ser Leu Phe Gln Arg Asp Ser Glu Thr Lys Gly Glu Ser
                155
                                     160
                                                          165
Gly Leu Val Leu Glu Gly Asp Lys Glu Ile His Gln Ile Phe Glu
                170
                                     175
                                                          180
Asp Leu Asp Lys Lys Leu Ala Leu Ala Ser Arg Phe Tyr Ile Pro
                185
                                     190
Glu Gly Cys Ile Gln Arg Trp Ala Ala Glu Met Val Val Ala Leu
                200
                                     205
                                                          210
Asp Ala Leu His Arg Glu Gly Ile Val Cys Arg Asp Leu Asn Pro
                215
                                     220
Asn Asn Ile Leu Leu Asn Asp Arg Gly His Ile Gln Leu Thr Tyr
                230
                                     235
Phe Ser Arg Trp Ser Glu Val Glu Asp Ser Cys Asp Ser Asp Ala
                245
                                     250
Ile Glu Arg Met Tyr Cys Ala Pro Glu Val Gly Ala Ile Thr Glu
                260
                                     265
                                                         270
Glu Thr Glu Ala Cys Asp Trp Trp Ser Leu Gly Ala Val Leu Phe
                275
                                     280
                                                         285
Glu Leu Leu Thr Gly Lys Thr Leu Val Glu Cys His Pro Ala Gly
                290
                                    295
                                                         300
Ile Asn Thr His Thr Thr Leu Asn Met Pro Glu Cys Val Ser Glu
                305
                                    310
                                                         315
Glu Ala Arg Ser Leu Ile Gln Gln Leu Leu Gln Phe Asn Pro Leu
                320
                                    325
Glu Arg Leu Gly Ala Gly Val Ala Gly Val Glu Asp Ile Lys Ser
                335
                                     340
His Pro Phe Phe Thr Pro Val Asp Trp Ala Glu Leu Met Arg
                350
                                    355
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<210> 53 <211> 545

<210> 52

<212> PRT <213> Homo sapiens <220> <221> misc feature <223> Incyte clone 1645941CD1 <400> 53 Met Ser Arg Lys Gln Asn Gln Lys Asp Ser Ser Gly Phe Ile Phe Asp Leu Gln Ser Asn Thr Val Leu Ala Gln Gly Gly Ala Phe Glu Asn Met Lys Glu Lys Ile Asn Ala Val Arg Ala Ile Val Pro Asn Lys Ser Asn Asn Glu Ile Ile Leu Val Leu Gln His Phe Asp Asn Cys Val Asp Lys Thr Val Gln Ala Phe Met Glu Gly Ser Ala Ser Glu Val Leu Lys Glu Trp Thr Val Thr Gly Lys Lys Lys Asn Lys Lys Lys Lys Asn Lys Pro Lys Pro Ala Ala Glu Pro Ser Asn Gly Ile Pro Asp Ser Ser Lys Ser Val Ser Ile Gln Glu Glu Gln Ser Ala Pro Ser Ser Glu Lys Gly Gly Met Asn Gly Tyr His Val Asn Gly Ala Ile Asn Asp Thr Glu Ser Val Asp Ser Leu Ser Glu Gly Leu Glu Thr Leu Ser Ile Asp Ala Arg Glu Leu Glu Asp Pro Glu Ser Ala Met Leu Asp Thr Leu Asp Arg Thr Gly Ser Met Leu Gln Asn Gly Val Ser Asp Phe Glu Thr Lys Ser Leu Thr Met His Ser Ile His Asn Ser Gln Gln Pro Arg Asn Ala Ala Lys Ser Leu Ser Arg Pro Thr Thr Glu Thr Gln Phe Ser Asn Met Gly Met Glu Asp Val Pro Leu Ala Thr Ser Lys Lys Leu Ser Ser Asn Ile Glu Lys Ser Val Lys Asp Leu Gln Arg Cys Thr Val Ser Leu Ala Arg Tyr Arg Val Val Lys Glu Glu Met Asp Ala Ser Ile Lys Lys Met Lys Gln Ala Phe Ala Glu Leu Glu Ser Cys Leu Met Asp Arg Glu Val Ala Leu Leu Ala Glu Met Asp Lys Val Lys Ala Glu Ala Met Glu Ile Leu Leu Ser Arg Gln Lys Lys Ala Glu Leu Leu Lys Lys Met Thr His Val Ala Val Gln Met Ser Glu Gln Gln Leu Val Glu Leu Arg Ala Asp Ile Lys His Phe Val Ser Glu Arg Lys Tyr Asp Glu Asp Leu Gly Arg Val Ala Arg Phe Thr Cys Asp Val Glu Thr Leu Lys Lys Ser Ile Asp Ser Phe Gly Gln Val Ser His Pro Lys Asn Ser Tyr Ser Thr Arg Ser Arg Cys Ser Ser Val Thr Ser Val 

Ser Leu Ser Ser Pro Ser Asp Ala Ser Ala Ala Ser Ser Ser Thr

Cys Ala Ser Pro Pro Ser Leu Thr Ser Ala Asn Lys Lys Asn Phe

```
Ala Pro Gly Glu Thr Pro Ala Ala Ile Ala Asn Ser Ser Gly Gln
                 425
                                      430
                                                          435
Pro Tyr Gln Pro Leu Arg Glu Val Leu Pro Gly Asn Arg Arg Gly
                 440
                                      445
                                                          450
Gly Gln Gly Tyr Arg Pro Gln Gly Gln Lys Ser Asn Asp Pro Met
                 455
                                      460
                                                          465
Asn Gln Gly Arg His Asp Ser Met Gly Arg Tyr Arg Asn Ser Ser
                 470
                                      475
Trp Tyr Ser Ser Gly Ser Arg Tyr Gln Ser Ala Pro Ser Gln Ala
                 485
                                     490
Pro Gly Asn Thr Ile Glu Arg Gly Gln Thr His Ser Ala Gly Thr
                 500
                                     505
Asn Gly Thr Gly Val Ser Met Glu Pro Ser Pro Pro Thr Pro Ser
                 515
                                     520
                                                          525
Phe Lys Lys Gly Leu Pro Gln Arg Lys Pro Arg Thr Ser Gln Thr
                 530
                                     535
Glu Ala Val Asn Ser
                 545
<210> 54
<211> 99
<212> PRT
<213> Homo sapiens
<220>
<221> misc feature
<223> Incyte clone 1646005CD1
<400> 54
Met Asn Trp Val Ala Val Leu Cys Pro Leu Gly Ile Val Trp Met
                                      10
                                                           15
Val Gly Asp Gln Pro Pro Gln Val Leu Ser Gln Ala Ser Ser Leu
                 20
                                      25
                                                           30
Ala Val Tyr Leu Arg Ala Ala Pro Tyr Pro Asp Val Thr Ala Lys
                 35
                                      40
                                                           45
Lys Leu Arg His Asp Thr Asn Cys Gly Phe Pro Arg Gln Gln Arg
                                      55
                                                           60
Met Ala Arg Gly His Glu Gly Arg Ala Pro Leu Leu Asp Arg Pro
                 65
                                                           75
Thr Leu Lys Ser Arg Tyr Leu Arg Ala Asn His Lys Ile Asn Thr
                 80
                                                           90
Phe Glu Glu Ile Thr Ala Met Pro Ser
                 95
<210> 55
<211> 565
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 1686561CD1
<400> 55
Met Asn Arg Ser Ile Pro Val Glu Val Asp Glu Ser Glu Pro Tyr
 1
                                      10
                                                           15
Pro Ser Gln Leu Lys Pro Ile Pro Glu Tyr Ser Pro Glu Glu
                 20
                                      25
Glu Ser Glu Pro Pro Ala Pro Asn Ile Arg Asn Met Ala Pro Asn
                 35
                                      40
                                                           45
Ser Leu Ser Ala Pro Thr Met Leu His Asn Ser Ser Gly Asp Phe
                 50
```

55

Ser	Gln	Ala	His	Ser 65	Thr	Leu	Lys	Leu	Ala 70	Asn	His	Gln	Arg	Pro 75
Val	Ser	Arg	Gln	Val 80	Thr	Cys	Leu	Arg	Thr 85	Gln	Val	Leu	Glu	
				95			Arg		100				_	Ala 105
Phe	Pro	Ser	Gly	Cys 110	Ser	Ala	Val	Ser	Glu 115	Pro	Ala	Ser	Glu	Ser 120
Val	Val	Gly	Ala	Leu 125	Pro	Ala	Glu	His	Gln 130	Phe	Ser	Phe	Met	
				140			Ser		145					Pro 150
				155			Lys		160					165
				170			Gly		175					Arg 180
				185			Ala		190					Ile 195
				200			Pro		205					210
				215			Leu		220					225
				230			Ser		235					240
				245			Ala		250					255
				260			Tyr		265					270
				275			Gly		280					285
				2 <del>9</del> 0			Ala		295					300
				305			His		310		_			315
				320			Gln		325					33Ō
				335			Pro		340					345
				350			Gly		355					360
				365			Gln		370					375
				380			Ser		385			_	_	390
				395			Glu		400					405
				410			Met		415		-			420
				425			Gln		430					435
				440			Ile		445					450
				455			Ile		460					465
				470			Ala		475					480
				485			Tyr		490					495
				500			Met		505					510
				515			Glu		520			-		525
Asn	Thr	His	Val	Tyr 530	Ser	Trp	Pro	Lys	Asn 535	Lys	Lys	Asn	Ile	Leu 540

Leu Arg Leu Leu Arg Glu Glu Glu Tyr Val Ala Pro Pro Arg Gly
545 . 550 . 555

Pro Leu Pro Thr Leu Gln Val Val Pro Leu
560 . 565

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<210> 56
<211> 197
<212> PRT
<213> Homo sapiens
<220>
<221> misc feature
<223> Incyte clone 1821233CD1
<400> 56
Met Thr Pro Thr Ser Ser Phe Val Ser Pro Pro Pro Pro Thr Ala
                                      10
Ser Pro His Ser Asn Arg Thr Thr Pro Pro Glu Ala Ala Gln Asn
                 20
                                      25
Gly Gln Ser Pro Met Ala Ala Leu Ile Leu Val Ala Asp Asn Ala
                                      40
                                                           45
Gly Gly Ser His Ala Ser Lys Asp Ala Asn Gln Val His Ser Thr
                 50
                                                           60
Thr Arg Arg Asn Ser Asn Ser Pro Pro Ser Pro Ser Ser Met Asn
                 65
                                      70
Gln Arg Arg Leu Gly Pro Arg Glu Val Gly Gly Gln Gly Ala Gly
                 80
                                      85
Asn Thr Gly Gly Leu Glu Pro Val His Pro Ala Ser Leu Pro Asp
                 95
                                     100
                                                         105
Ser Ser Leu Ala Thr Ser Ala Pro Leu Cys Cys Thr Leu Cys His
                110
                                     115
                                                         120
Glu Arg Leu Glu Asp Thr His Phe Val Gln Cys Pro Ser Val Pro
                125
                                     130
                                                         135
Ser His Lys Phe Cys Phe Pro Cys Ser Arg Gln Ser Ile Lys Gln
                140
                                     145
Gln Gly Ala Ser Gly Glu Val Tyr Cys Pro Ser Gly Glu Lys Cys
                155
                                     160
                                                         165
Pro Leu Val Gly Ser Asn Val Pro Trp Ala Phe Met Gln Gly Glu
                170
                                     175
                                                         180
Ile Ala Thr Ile Leu Ala Gly Asp Val Lys Val Lys Glu Arg
                                     190
                                                         195
Asp Ser
```

<210> 57 <211> 321 <212> PRT

<213> Homo sapiens

<220>

<221> misc\_feature

<223> Incyte clone 1877278CD1

<400> 57

Met Lys Glu Asp Cys Leu Pro Ser Ser His Val Pro Ile Ser Asp 1 5 10 15 Ser Lys Ser Ile Gln Lys Ser Glu Leu Leu Gly Leu Leu Lys Thr 20 25 30 Tyr Asn Cys Tyr His Glu Gly Lys Ser Phe Gln Leu Arg His Arg 35 40 45

```
Glu Glu Glu Gly Thr Leu Ile Ile Glu Gly Leu Leu Asn Ile Ala
                 50
                                      55
Trp Gly Leu Arg Arg Pro Ile Arg Leu Gln Met Gln Asp Asp Arg
                 65
                                      70
                                                          75
Glu Gln Val His Leu Pro Ser Thr Ser Trp Met Pro Arg Arg Pro
                                                          90
                 80
                                      85
Ser Cys Pro Leu Lys Glu Pro Ser Pro Gln Asn Gly Asn Ile Thr
                 95
                                     100
Ala Gln Gly Pro Ser Ile Gln Pro Val His Lys Ala Glu Ser Ser
                110
                                    115
Thr Asp Ser Ser Gly Pro Leu Glu Glu Ala Glu Glu Ala Pro Gln
                125
                                    130
Leu Met Arg Thr Lys Ser Asp Ala Ser Cys Met Ser Gln Arg Arg
                140
                                    145
                                                         150
Pro Lys Cys Arg Ala Pro Gly Glu Ala Gln Arg Ile Arg Arg His
                155
                                    160
                                                         165
Arg Phe Ser Ile Asn Gly His Phe Tyr Asn His Lys Thr Ser Val
                170
                                    175
                                                         180
Phe Thr Pro Ala Tyr Gly Ser Val Thr Asn Val Arg Val Asn Ser
                185
                                    190
                                                         195
Thr Met Thr Thr Leu Gln Val Leu Thr Leu Leu Asn Lys Phe
                200
                                     205
Arg Val Glu Asp Gly Pro Ser Glu Phe Ala Leu Tyr Ile Val His
                215
                                     220
                                                         225
Glu Ser Gly Glu Arg Thr Lys Leu Lys Asp Cys Glu Tyr Pro Leu
                230
                                     235
Ile Ser Arg Ile Leu His Gly Pro Cys Glu Lys Ile Ala Arg Ile
                245
                                     250
Phe Leu Met Glu Ala Asp Leu Gly Val Glu Val Pro His Glu Val
                260
                                     265
                                                         270
Ala Gln Tyr Ile Lys Phe Glu Met Pro Val Leu Asp Ser Phe Val
                275
                                    280
                                                         285
Glu Lys Leu Lys Glu Glu Glu Glu Arg Glu Ile Ile Lys Leu Thr
                290
                                    295
                                                         300
Met Lys Phe Gln Ala Leu Arg Leu Thr Met Leu Gln Arg Leu Glu
                305
                                    310
                                                         315
Gln Leu Val Glu Ala Lys
                320
```

```
<210> 58
<211> 356
<212> PRT
<213> Homo sapiens
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<220>

<221> misc\_feature

<223> Incyte clone 1880692CD1

<400> 58 Met Glu Trp Leu Lys Ser Thr Asp Tyr Gly Lys Tyr Glu Gly Leu Thr Lys Asn Tyr Met Asp Tyr Leu Ser Arg Leu Tyr Glu Arg Glu Ile Lys Asp Phe Phe Glu Val Ala Lys Ile Lys Met Thr Gly Thr Thr Lys Glu Ser Lys Lys Phe Gly Leu His Gly Ser Ser Gly Lys Leu Thr Gly Ser Thr Ser Ser Leu Asn Lys Leu Ser Val Gln Ser Ser Gly Asn Arg Arg Ser Gln Ser Ser Leu Leu Asp Met Gly Asn Met Ser Ala Ser Asp Leu Asp Val Ala Asp Arg Thr Lys Phe 

```
Asp Lys Ile Phe Glu Gln Val Leu Ser Glu Leu Glu Pro Leu Cys
                110
                                     115
                                                          120
Leu Ala Glu Gln Asp Phe Ile Ser Lys Phe Phe Lys Leu Gln Gln
                125
                                     130
                                                          135
His Gln Ser Met Pro Gly Thr Met Ala Glu Ala Glu Asp Leu Asp
                140
                                     145
                                                          150
Gly Gly Thr Leu Ser Arg Gln His Asn Cys Gly Thr Pro Leu Pro
                155
                                     160
Val Ser Ser Glu Lys Asp Met Ile Arg Gln Met Met Ile Lys Ile
                170
                                     175
Phe Arg Cys Ile Glu Pro Glu Leu Asn Asn Leu Ile Ala Leu Gly
                185
                                     190
Asp Lys Ile Asp Ser Phe Asn Ser Leu Tyr Met Leu Val Lys Met
                200
                                     205
Ser His His Val Trp Thr Ala Gln Asn Val Asp Pro Ala Ser Phe
                215
                                     220
                                                          225
Leu Ser Thr Thr Leu Gly Asn Val Leu Val Thr Val Lys Arg Asn
                230
                                     235
Phe Asp Lys Cys Ile Ser Asn Gln Ile Arg Gln Met Glu Glu Val
                245
                                     250
                                                          255
Lys Ile Ser Lys Lys Ser Lys Val Gly Ile Leu Pro Phe Val Ala
                260
                                     265
Glu Phe Glu Glu Phe Ala Gly Leu Ala Glu Ser Ile Phe Lys Asn
                275
                                     280
                                                          285
Ala Glu Arg Arg Gly Asp Leu Asp Lys Ala Tyr Thr Lys Leu Ile
                290
                                     295
Arg Gly Val Phe Val Asn Val Glu Lys Val Ala Asn Glu Ser Gln
                305
                                     310
                                                          315
Lys Thr Pro Arg Asp Val Val Met Met Glu Asn Phe His His Ile
                320
                                     325
Phe Ala Thr Leu Ser Arg Leu Lys Ile Ser Cys Leu Glu Ala Glu
                335
                                     340
Lys Lys Glu Ala Ala Ile Asn His Lys Phe Phe
                350
                                     355
```

<210> 59 <211> 299 <212> PRT <213> Homo sapiens <220>

<221> misc\_feature <223> Incyte clone 2280456CD1

<400> 59 Met Glu Glu Leu Leu Pro Asp Gly Gln Ile Trp Ala Asn Met Asp Pro Glu Glu Arg Met Leu Ala Ala Ala Thr Ala Phe Thr His Ile Cys Ala Gly Gln Gly Glu Gly Asp Val Arg Arg Glu Ala Gln Ser Ile Gln Tyr Asp Pro Tyr Ser Lys Ala Ser Val Ala Pro Gly Lys Arg Pro Ala Leu Pro Val Gln Leu Gln Tyr Pro His Val Glu Ser Asn Val Pro Ser Glu Thr Val Ser Glu Ala Ser Gln Arg Leu Arg Lys Pro Val Met Lys Arg Lys Val Leu Arg Arg Lys Pro Asp Gly Glu Val Leu Val Thr Asp Glu Ser Ile Ile Ser Glu Ser Glu Ser Gly Thr Glu Asn Asp Gln Asp Leu Trp Asp Leu Arg Gln Arg Leu 

```
Met Asn Val Gln Phe Gln Glu Asp Lys Glu Ser Ser Phe Asp Val
                140
                                     145
                                                          150
Ser Gln Lys Phe Asn Leu Pro His Glu Tyr Gln Gly Ile Ser Gln
                155
                                     160
Asp Gln Leu Ile Cys Ser Leu Gln Arg Glu Gly Met Gly Ser Pro
                170
                                     175
                                                          180
Ala Tyr Glu Gln Asp Leu Ile Val Ala Ser Arg Pro Lys Ser Phe
                185
                                     190
                                                          195
Ile Leu Pro Lys Leu Asp Gln Leu Ser Arg Asn Arg Gly Lys Thr
                200
                                     205
                                                          210
Asp Arg Val Ala Arg Tyr Phe Glu Tyr Lys Arg Asp Trp Asp Ser
                215
                                     220
Ile Arg Leu Pro Gly Glu Asp His Arg Lys Glu Leu Arg Trp Gly
                230
                                     235
Val Arg Glu Gln Met Leu Cys Arg Ala Glu Pro Gln Ser Lys Pro
                245
                                     250
                                                          255
Gln His Ile Tyr Val Pro Asn Asn Tyr Leu Val Pro Thr Glu Lys
                260
                                     265
                                                          270
Lvs Arg Ser Ala Leu Arg Trp Gly Val Arg Cys Asp Leu Ala Asn
                275
                                     280
                                                          285
Gly Val Ile Pro Arg Lys Leu Pro Phe Pro Leu Ser Pro Ser
                290
                                     295
```

<210> 60 <211> 293 <212> PRT <213> Homo sapiens <220> <221> misc\_feature <223> Incyte clone 2284580CD1 <400> 60 Met Ala Thr Phe Ser Gly Pro Ala Gly Pro Ile Leu Ser Leu Asn Pro Gln Glu Asp Val Glu Phe Gln Lys Glu Val Ala Gln Val Arg Lys Arg Ile Thr Gln Arg Lys Lys Gln Glu Gln Leu Thr Pro Gly Val Val Tyr Val Arg His Leu Pro Asn Leu Leu Asp Glu Thr Gln Ile Phe Ser Tyr Phe Ser Gln Phe Gly Thr Val Thr Arg Phe Arg Leu Ser Arg Ser Lys Arg Thr Gly Asn Ser Lys Gly Tyr Ala Phe Val Glu Phe Glu Ser Glu Asp Val Ala Lys Ile Val Ala Glu Thr Met Asn Asn Tyr Leu Phe Gly Glu Arg Leu Leu Glu Cys His Phe Met Pro Pro Glu Lys Val His Lys Glu Leu Phe Lys Asp Trp Asn Ile Pro Phe Lys Gln Pro Ser Tyr Pro Ser Val Lys Arg Tyr Asn Arg Asn Arg Thr Leu Thr Gln Lys Leu Arg Met Glu Glu Arg Phe Lys Lys Glu Arg Leu Leu Arg Lys Lys Leu Ala Lys Lys Gly Ile Asp Tyr Asp Phe Pro Ser Leu Ile Leu Gln Lys Thr Glu Ser Ile Ser Lys Thr Asn Arg Gln Thr Ser Thr Lys Gly Gln Val Leu 

```
Arg Lys Lys Lys Lys Val Ser Gly Thr Leu Asp Thr Pro Glu
                215
                                     220
Lys Thr Val Asp Ser Gln Gly Pro Thr Pro Val Cys Thr Pro Thr
                230
                                     235
                                                          240
Phe Leu Glu Arg Arg Lys Ser Gln Val Ala Glu Leu Asn Asp Asp
                245
                                     250
                                                          255
Asp Lys Asp Asp Glu Ile Val Phe Lys Gln Pro Ile Ser Cys Val
                260
                                     265
                                                          270
Lys Glu Glu Ile Gln Glu Thr Gln Thr Pro Thr His Ser Arg Lys
                275
                                                          285
Lys Arg Arg Arg Ser Ser Asn Gln
                290
```

<210> 61

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<211> 777
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<223> Incyte clone 2779172CD1
<400> 61
Met Val Leu Cys His Ser Phe Leu Tyr Arg Ile Leu Thr Val Gln
                                      10
                                                           15
Gln His Gly Phe Phe Gly His Asp Arg Arg Pro Ala Asp Gly
                  20
                                      25
                                                           30
Glu Lys Gln Ala Ala Thr His Val Ser Leu Asp Gln Glu Tyr Asp
                  35
                                      40
                                                           45
Ser Glu Ser Ser Gln Gln Trp Arg Glu Leu Glu Glu Gln Val Val
                 50
                                      55
                                                           60
Ser Val Val Asn Lys Gly Val Ile Pro Ser Asn Phe His Pro Thr
                 65
                                      70
                                                           75
Gln Tyr Cys Leu Asn Ser Tyr Ser Asp Asn Ser Arg Phe Pro Leu
                 80
                                      8.5
                                                           90
Ala Val Val Glu Glu Pro Ile Thr Val Glu Val Ala Phe Arg Asn
                 95
                                     100
                                                          105
Pro Leu Lys Val Leu Leu Leu Thr Asp Leu Ser Leu Leu Trp
                110
                                     115
                                                          120
Lys Phe His Pro Lys Asp Phe Ser Gly Lys Asp Asn Glu Glu Val
                125
                                     130
Lys Gln Leu Val Thr Ser Glu Pro Glu Met Ile Gly Ala Glu Val
                140
                                     145
Ile Ser Glu Phe Leu Ile Asn Gly Glu Glu Ser Lys Val Ala Arg
                155
                                     160
                                                          165
Leu Lys Leu Phe Pro His His Ile Gly Glu Leu His Ile Leu Gly
                170
                                     175
                                                          180
Val Val Tyr Asn Leu Gly Thr Ile Gln Gly Ser Met Thr Val Asp
                185
                                     190
                                                          195
Gly Ile Gly Ala Leu Pro Gly Cys His Thr Gly Lys Tyr Ser Leu
                200
                                     205
                                                          210
Ser Met Ser Val Arg Gly Lys Gln Asp Leu Glu Ile Gln Gly Pro
                215
                                     220
                                                          225
Arg Leu Asn Asn Thr Lys Glu Glu Lys Thr Ser Val Lys Tyr Gly
                230
                                     235
                                                          240
Pro Asp Arg Arg Leu Asp Pro Ile Ile Thr Glu Glu Met Pro Leu
                245
                                     250
                                                          255
Leu Glu Val Phe Phe Ile His Phe Pro Thr Gly Leu Leu Cys Gly
                260
                                     265
                                                          270
Glu Ile Arg Lys Ala Tyr Val Glu Phe Val Asn Val Ser Lys Cys
                275
                                     280
                                                          285
```

Pro	Leu	Thr	Gly	Leu 290	Lys	Val	Val	Ser	Lys 295	Arg	Pro	Glu	Phe	Phe 300
Thr	Phe	Gly	Gly		Thr	Ala	Val	Leu		Pro	Leu	Ser	Pro	
Ala	Ser	Glu	Asn		Ser	Ala	Tyr	Lys		Val	Val	Thr	Asp	
Thr	Ser	Val	Cys	Thr 335	Ala	Leu	Ile	Ser		Ala	Ser	Ser	Val	
Phe	Gly	Ile	Gly	Thr 350	Gly	Ser	Gln	Pro		Val	Ile	Pro	Val	
Leu	Pro	Asp	Thr	Val 365	Leu	Leu	Pro	Gly	Ala 370	Ser	Val	Gln	Leu	
Met	Trp	Leu	Arg	Gly 380	Pro	Asp	Glu	Glu	Gly 385	Val	His	Glu	Ile	Asn 390
				395					400		Pro	_		405
				410					415		Thr			420
				425					430		Ser			435
				440					445		Val	_		450
				455					460		Phe			465
				470					475		Gln			480
				485					490		Ser			495
				500					505		Lys			510
				515					520		Asp			525
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				560					565		Asp			570
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				695					700		Glu			705
				710					715		Gln			720
				725					730		Leu			735
				740					745		Pro			750
Ala	Lys	Leu	Ser	Asp 755	Gln	Val	Thr	Val	Phe 760	Glu	Thr	Ser	Gln	Gln 765

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Leu Leu Cys Phe Trp Val Asn Arg Cys Ala Cys Gln Leu Ala
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                                      40
                                                          45
Cys Val Leu Ser Lys Phe His Lys Leu Lys Val Phe Lys Gly Cys
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Val Val Ser Glu Leu Tyr Val Ser Phe Leu Ser Leu Tyr Leu Gln
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                                      40
Pro His Ser Leu Pro Pro Pro Leu Thr Leu His Cys Asn Ile Thr
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Glu Ser Tyr Pro Ser Ser Ser Pro Ile Trp Phe Val Asp Ser Glu
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Asp Pro Asn Leu Thr Ser Val Leu Glu Arg Leu Glu Asp Thr Lys
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Asn Asn Asn Leu Asn Gly Thr Thr Glu Glu Val Thr Ser Glu Glu
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Glu Glu Glu Glu Glu Met Ala Glu Asp Ile Glu Asp Leu Asp
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His Tyr Glu Met Lys Glu Glu Glu Pro Ile Ser Gly Lys Lys Ser
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                                     130
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Glu Asp Glu Gly Ile Glu Lys Glu Asn Leu Ala Ile Leu Glu Lys
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Leu Ile Asn Asp Ser Leu Tyr Asp Trp His Val Lys Leu Gln Lys
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Val Asp Pro Asp Ser Pro Leu His Ser Asp Leu Gln Ile Leu Lys
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Glu Lys Glu Gly Ile Glu Tyr Ile Leu Leu Asn Phe Ser Phe Lys
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Asp Asn Phe Pro Phe Asp Pro Pro Phe Val Arg Val Leu Pro
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Gln Ser Tyr Asn Ser Ile Val Gln Ile His Glu Lys Asn Gly Trp
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Tyr Thr Pro Pro Lys Glu Asp Gly
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Gly Gly Ser Leu Ile Ala Glu Gln Trp Val Leu Thr Ala Ala His
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Cys Phe Arg Asn Thr Ser Glu Thr Ser Leu Tyr Gln Val Leu Leu
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Gly Ala Arg Gln Leu Val Gln Pro Gly Pro His Ala Met Tyr Ala
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Arg Val Arg Gln Val Glu Ser Asn Pro Leu Tyr Gln Gly Thr Ala
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Ser Ser Ala Asp Val Ala Leu Val Glu Leu Glu Ala Pro Val Pro
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Phe Thr Asn Tyr Ile Leu Pro Val Cys Leu Pro Asp Pro Ser Val
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Ile Phe Glu Thr Gly Met Asn Cys Trp Val Thr Gly Trp Gly Ser
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                                                         165
Pro Ser Glu Glu Asp Leu Leu Pro Glu Pro Arg Ile Leu Gln Lys
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Leu Ala Val Pro Ile Ile Asp Thr Pro Lys Cys Asn Leu Leu Tyr
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                                                         195
Ser Lys Asp Thr Glu Phe Gly Tyr Gln Pro Lys Thr Ile Lys Asn
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Asp Met Leu Cys Ala Gly Phe Glu Glu Gly Lys Lys Asp Ala Cys
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Lys Gly Asp Ser Gly Gly Pro Leu Val Cys Leu Val Gly Gln Ser

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                                                          135
Val Asp Phe Arg Arg Leu Gly Glu Glu Phe Cys His Trp Phe Phe
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Glu Trp Gly Pro Gln His Phe Trp His Asp Val Lys Leu Arg Phe
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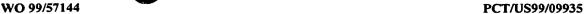


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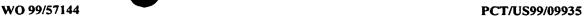
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WO 99/57144

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<223> Incyte clone 1645941CB1

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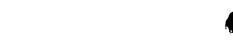
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WO 99/57144

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